PCT

WORLD INTELLECTUAL PROPERTY ORGANIZATION International Bureau



INTERNATIONAL APPLICATION PUBLISHED UNDER THE PATENT COOPERATION TREATY (PCT)

(51) International Patent Classificati n ⁵:

C12N 15/11, C12Q 1/68

(11) International Publication Number:

WO 93/16178

A2

(43) International Publication Date:

19 August 1993 (19.08.93)

(21) International Application Number:

PCT/US93/01294

(22) International Filing Date:

12 February 1993 (12.02.93)

(30) Priority data:

07/837,195

12 February 1992 (12.02.92) US

rualy 1992 (12.02.92) OS

(71) Applicant: THE UNITED STATES OF AMERICA, as represented by THE SECRETARY, DEPARTMENT OF HEALTH AND HUMAN SERVICES [US/US]; Washington, DC (US).

(72) Inventors: VENTER, Craig, J.; 1718 Nordic Hill Circle, Silver Spring, MD 20906 (US). ADAMS, Mark, D.; 12812 Sage Terrace, Germantown, MD 20874 (US). MORENO, Ruben, F.; 14415 Coral Gables Way, North Potomac, MD 20878 (US).

(74) Agents: ALTMAN, Daniel, E. et al.; Knobbe, Martens, Olson and Bear, 620 Newport Center Drive, 16th Floor, Newport Beach, CA 92660 (US).

(81) Designated States: AU, CA, JP, European patent (AT, BE, CH, DE, DK, ES, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE).

Published

Without international search report and to be republished upon receipt of that report.

(54) Title: SEQUENCES CHARACTERISTIC OF HUMAN GENE TRANSCRIPTION PRODUCT

(57) Abstract

Partial and complete human cDNA and genomic sequences corresponding to particular expressed sequence tags (ESTs). The ESTs are cDNA sequences that are generally between 150 and 500 base pairs in length, are derived from human brain cDNA libraries, correspond to genes transcribed in human brain, and have base sequences identified herein as SEQ ID NOS: 1-2421.

FOR THE PURPOSES OF INFORMATION ONLY

Codes used to identify States party to the PCT on the front pages of pamphlets publishing international applications under the PCT.

AT	Austria	FR	France	MR	Mauritania
AU	Australia	GA	Gabon	MW	Malawi
88	Barbados	CB	United Kingdom	NL	Netherlands
BE	Belgium	· GN	Guinea	NO	Norway
BF	Burkina Faso	CR	Greece	NZ	New Zealand
BG	Bulgaria	Hυ	Hungary	PL	Poland
BJ	Benin	16	Ireland	PT	Portugal
BR	Brazil	IT	Italy	RO	Romania
CA	Canada	JP	Japan	RU	Russian Federation
CF	Central African Republic	KP	Democratic People's Republic	SD	Sudan
CG	Congo		of Korea	SE	Sweden
СН	Switzerland	KR	Republic of Korea	SK	Slovak Republic
CI	Côte d'Ivoire	KZ	Kazakhstan	ŠN	Senegal
CM	Cameroon		Liechtenstein	SU	Soviet Union
CS	Czechoslovakia	LK	Sri Lanka	TD	Chad
CZ	Czech Republic	1.ti	Lasembourg	TG	Togo
DE	Germany	MC	Monaco	UA	Ukraine
DK	Denmark	MG	Madagascar	US	United States of America
ES	Spain	Ml.	Mali	VN	Viet Nam
FI	Finland	MN	Moneolia	***	

SEQUENCES CHARACTERISTIC OF HUMAN GENE TRANSCRIPTION PRODUCT

5

10

15

20

25

30

Technical Field

The present invention relates to newly identified polynucleotide sequences corresponding to transcription products of human genes, and to complete gene sequences associated therewith.

Background

This invention relates to human genes. Identification and sequencing of human genes is a major goal of modern scientific research. The sequence of human genes is more than just a scientific curiosity. For example, identifying genes and determining their sequences, scientists have been able to make large quantities of valuable human "gene products." These include human insulin, interferon, Factor VIII, tumor necrosis factor, human growth hormone, tissue plasminogen activator, and numerous other compounds. Additionally, knowledge of gene sequences can provide the k y to treatment or cure of genetic diseases (such as muscular dystrophy and cystic fibrosis). The present invention represents a quantum leap forward in mankind's knowledge of human gene sequences.

There are several basic concepts of molecular biology which figure prominently in the invention. A brief explanation of those concepts follows. Additional background information and definitions for scientific terms can be found

10

15

20

25

30

35

in the literature. See, for example, "Glossary of Genetics, Classical and Molecular" by R. Rieger, A. Michaelis, and M.M. Green (Fifth Edition, Springer-Verlag, New York (1991)). The contents of this and other publications cited in the specification are incorporated by reference herein.

At an initial level, the present invention is based on identification and characterization of gene segments. Genes are the basic units of inheritance. Each gene is a string of connected bases called nucleotides. Most genes are formed of deoxyribonucleic acid, DNA. (Some viruses contain genes of ribonucleic acid, RNA.) The genetic information resides in the particular sequence in which the bases are arranged. A short sequence of nucleotides is often called a polynucleotide or an oligonucleotide.

Like genes, polypeptides are built from long strings of individual units. These units are amino acids. nucleotide sequence of a gene tells the cell the sequence in which to arrange the amino acids to make the polypeptide encoded by that gene. In general, chains of up to about 200 amino acids are called polypeptides, while proteins are larger molecules made up of polypeptide subunits; both types of molecules are referred to generally herein as polypeptides. A triplet of nucleotides (codon) in DNA codes for each amino acid or signals the beginning or end of the message (anticodon). The term codon is also used for the corresponding (and complementary) sequences of three nucleotides in the mRNA into which the original DNA sequence is transcribed.

Generally, enzymes in the cell transcribe the permanent DNA of the gene into a temporary RNA copy, called messenger RNA or mRNA. The mRNA, in turn, can be translated into a polypeptide by the cell. This entire process is called gene expression, and the polypeptide is the gene product encoded by the gene.

Scientists have previously discovered how to reverse the transcription process and copy mRNA back into DNA using an

-3-

enzyme called reverse transcriptase. The resulting is called complementary DNA, or cDNA. This is schematically shown in the single Figure. When substantially all of the mRNA from one cell or tissue is converted to cDNA at once and cloned into multiple copies of a recombinant vector to allow replication and manipulation in the laboratory, the result is called a cDNA library.

5

10

15

20

25

. 30

35

The various types of genes include those which code for polypeptides, those which are transcribed into RNA but are not translated into polypeptides, and those whose functional significance does not demand that they be transcribed at all. Most genes are found on large molecules of DNA located in chromosomes. Double stranded CDNA carries all information of a gene. Each base of the first strand is joined to a complementary base (hybridized) in the second linear DNA molecules in chromosomes have strand. The thousands of genes distributed along their Chromosomes include both coding regions (coding polypeptides) and noncoding regions; the coding regions. represent only about three percent of the total chromosome sequence.

An individual gene has regulatory regions that include a promoter which directs expression of the gene, a coding region which can code for a polypeptide, and a termination signal. The regulatory DNA sequence is usually a noncoding region that determines if, where, when, and at what level a particular gene is expressed.

The coding regions of many genes are discontinuous, with coding sequences (exons) alternating with noncoding regions (introns). The final mRNA copy of the gene does not include these introns (which can be much longer than the coding region itself), although it does contain certain untranslated regions that usually do not code for the polynucleotide gene product. Untranslated sequences at the beginning and end of the mRNA are known as 5'- and 3'-untranslated regions,

15

20.

25

30

35

respectively. This nomenclature reflects the orientation of the nucleotide constituents of the mRNA.

A cDNA is a DNA copy of a messenger RNA, which contains all of the exons of a gene. The cDNA can be thought of as having three parts: an untranslated 5' leader, uninterrupted polypeptide-coding sequence, and 3 ' untranslated region. The untranslated leader and trailing sequences are important for initiation of translation, mRNA stability, and other functions. The untranslated leader and trailing sequences are called 5'- and 3'-untranslated sequences, respectively. The 3' untranslated sequence is usually longer than the 5' untranslated leader, and can be polypeptide-coding than the sequence. The regions typically untranslated have many, randomlydistributed stop codons, and do not display the nonrandom base arrangements found in coding sequences. untranslated sequence is relatively short, generally between 20 and 200 bases. The 3'-untranslated sequence is often many times longer, up to several thousand bases.

The translated or coding sequence begins with a translational start codon (AUG or GUG) and ends with a translational stop codon (UAA, UGA, or UAG). Generally, translation begins at the first "start" codon on the mRNA and proceeds to the first "stop" codon. Coding sequences can be distinguished by their nonrandom distribution of bases; numerous computer algorithms have been developed to distinguish coding from noncoding regions in this way.

Human DNA differs from person to person. No two persons (except perhaps identical twins) have identical DNA. While the differences, called allelic variations or polymorphisms, are slight on a molecular level, they account for most of the physical and other observable differences between individuals. It has been estimated that approximately 14 million sequence polymorphism differences exist between individuals.

10

15

20

25

3.0

35

The ability of one strand of DNA to attach or hybridize to a complementary strand has already been exploited for several purposes. For example, small pieces of DNA (15 to 25 base pairs long) can be made which will hybridize to longer strands of DNA which have a complementary sequence. short "primers" can be selected such that they hybridize to a specific, unique location on the longer strand. primers have hybridized to their target on the DNA, the polymerase chain reaction (PCR) can be employed to generate millions of copies of (or amplify) the particular segment of DNA between the locations to which two primers are bound. Briefly, this technique allows amplification of a DNA region situated between two convergent primers, oligonucleotide primers that hybridize to opposite strands. Primer extension proceeds inward across the region between the two primers, and the product of DNA synthesis of one primer serves as a template for the other primer. cycles of DNA denaturation, annealing of primers, extension result in an exponential increase in the number of copies of the region bounded by the primers.

Similarly, a labeled segment of single-stranded DNA can be hybridized to a longer DNA sequence, such as a chromosome, to mark a specific location on the longer sequence. Segments of DNA 50 bases long or longer that hybridize to a unique DNA location in the human genome are extremely unlikely to hybridize elsewhere in the human genome.

The Human Genome Project is an effort to sequence all human DNA (the human genome). The human genome is estimated to comprise 50,000 - 100,000 genes, up to 30,000 of which might be expressed in the brain (Sutcliffe, Ann. Rev. Neurosci. 11:157 (1988)). Once dedicated human chromosome sequencing begins in three to five years, it was expected that 12-15 years will be required to complete the sequence of the genome (Report of the Ad Hoc Program Advisory Committee on Complex Genomes, Reston, Va., Feb. 1988, D. Baltimore Ed. (NIH, Bethesda, Md, 1988)). At that rate, the majority of

human genes would remain unknown for at least the next decade. The present invention can greatly accelerate the pace at which human genes can be identified and mapped. Most gene researchers, in conjunction with publication of their results in this field, submit sequence data to the GenBank database. Prior to the present invention, GenBank listed the sequences of only a few thousand human genes and less than two hundred human brain mRNAs (GenBank Release 66.0, December, 1990).

10

15

20

The role of sequencing complementary DNA (cDNA), reverse transcribed from mRNA, as a part of the human genome project has been vigorously debated since the idea of determining the complete nucleotide sequence of humans first surfaced. The coding sequence of all human genes represents most of the information content of the genome, but only 3-5% of the total DNA. In contrast, cDNA (which is only made from the transcription product of active genes) is one-half to three-fourths (the remainder being 5'- and 3'-untranslated sequence) meaningful genetic information. Thus, some have argued that cDNA sequencing should take precedence over genomic sequencing (Brenner, CIBA Found. Symp. 149:6 (1990)). However, until now, such arguments have not been heeded.

. 25

3.0

35.

Genomic sequencing proponents have argued the difficulty of finding every mRNA expressed in all tissues, cell types, and developmental states, and that much valuable information from intronic and intergenic regions, including control and regulatory sequences, will be missed by cDNA sequencing. (Report of the Committee on Mapping and Sequencing the Human Genome, National Research Council (National Academy Press, Washington, D.C. 1988)). Further, sequencing of transcribed regions of the genome using cDNA libraries has heretofore been considered impractical or unsatisfactory. Libraries of cDNA were believed to be dominated by repetitive elements, mitochondrial genes, ribosomal RNA genes, and other nuclear genes comprising common or housekeeping sequences. It was believed that cDNA libraries would provide few sequences

-7-

corresponding to structural and regulatory polypeptides or peptides. See, for example, Putney, et al., Nature 302:718-721 (1983). Putney, et al. sequenced over 150 clones from a rabbit muscle cDNA library and identified clones for 13 of the 19 known muscle polypeptides, including one new isotype but no unknown coding sequences.

5

10

15

20

25

30

35

Another perceived drawback of cDNA sequencing was that some mRNAs are abundant, and some are rare. The cellular quantities of mRNA from various genes can vary by several orders of magnitude. This led critics to believe that most information obtained from cDNA sequencing would be repetitious and useless.

The present invention demonstrates that, despite such skepticism, cDNA sequencing now provides a rapid method for obtaining enormous amounts of valuable genetic information and DNA products of great utility for the biotechnology and pharmaceutical industries. Not only can many distinct cDNAs be isolated and sequenced, even partial cDNAs can be used, with conventional, well-understood methods, to isolate entire and to determine the chromosomal locations and biological functions of these genes. As is demonstrated here, fragments of only a few hundred bases are sufficient, in many cases, to identify the probable function of a new human gene if it is similar in structure to a gene from another animal, or from plants or bacteria. Similarly, even fragments of untranslated regions of a cDNA can be used to: i) isolate the coding sequence of the cDNA; ii) isolate the complete gene; iii) determine the position of the gene on a human chromosome, and hence the potential of the gene to cause a human genetic disease; and iv) determine the function of the gene by means of experiments in which the function of the native gene is disrupted by the addition of a short DNA fragment to the cell, e.g., using triple helix or antisense probes.

Because coding regions comprise such a small portion of the human genome, identification and mapping of transcribed

-8-

regions and coding regions of chromosomes is of significant interest. There is a corresponding need for reagents for identifying and marking coding regions and transcribed regions of chromosomes. Furthermore, such human sequences are valuable for chromosome mapping, human identification, identification of tissue type and origin, forensic identification, and locating disease-associated genes (i.e., genes that are associated with an inherited human disease, whether through mutation, deletion, faulty gene or expression) on the chromosome.

5

10

15

20

25

30

35

SUMMARY OF THE INVENTION

Contrary to the expectations of the scientific community, cDNA screening and sequencing techniques have now been used to discover a large number of heretofore unknown human genes. Disclosed herein are over 2,400 new human polynucleotide sequences. These sequences could represent up to 5% of all human genes. The novelty of these sequences has been established through comparison to both nucleotide sequence databases and amino acid sequence databases. Surprisingly, over 80% of the sequences generated were unrelated to any sequences previously described in the literature.

The sequences of the present invention were ascertained using a fast approach to cDNA characterization. This approach could facilitate the tagging of most expressed human genes within a few years at a fraction of the cost of complete genomic sequencing, provide new genetic markers, provide new DNA-based therapeutics and diagnostics, and provide other valuable nucleotide reagents.

The sequences disclosed herein, styled Expressed Sequence Tags ("ESTs"), are markers for human genes actually transcribed in vivo. Techniques are disclosed for using these ESTs to obtain the full coding region of the corresponding gene. The use of ESTs, complete coding sequences, or fragments thereof for marking chromosomes, for

10

15

20

25

30

35

mapping locations of expressed genes on chromosomes, for individual or forensic identification, for mapping locations of disease-associated genes, for identification of tissue type, and for preparation of antisense sequences, probes, and constructs is discussed in detail below. Unlike the random genomic DNA sequence tagged sites (STSs) (Olson et al., Science 245:1434 (1989)), ESTs point directly to expressed genes.

Various aspects of the present invention thus include the individual ESTs, corresponding partial and complete cDNA, genomic DNA, mRNA, antisense strands, triple helix probes, PCR primers, coding regions, and constructs. Also, where one skilled in the art is enabled by this specification to prepare expression vectors and polypeptide expression products, they are also within the scope of the present invention, along with antibodies, especially monoclonal antibodies, to such expression products.

BRIEF DESCRIPTION OF THE DRAWING

The single drawing Figure schematically illustrates the progression from chromosome to gene to mRNA to cDNA.

DETAILED DESCRIPTION OF THE INVENTION

The detailed description that follows provides not only the actual sequence of each new EST, but also explains how the ESTs were obtained, how to obtain the corresponding complete cDNA sequence and the corresponding genomic DNA sequence, how to make DNA constructs from the ESTs and corresponding sequences, how to use those sequences as reagents in molecular biology and other fields, how to produce gene products from the ESTs and corresponding sequences and antibodies to those gene products, and the functional categories of many ESTs and corresponding genes. Furthermore, numerous actual working examples and predictive

-10-

examples are provided to demonstrate and exemplify numerous aspects of the invention.

I. ESTs from cDNA Libraries

5

10

15

20

25

30

35

The sequences of the present invention were isolated from commercially available and custom made cDNA libraries using a rapid screening and sequencing technique. general, the method comprises applying conventional automated DNA sequencing technology to screening clones, advantageously randomly selected clones, from a cDNA library. Preferably, the library is initially "enriched" through removal of ribosomal sequences and other common sequences prior to clone selection. According to the present method, ESTs are generated from partial DNA sequencing of the selected clones. The ESTs of the present invention were generated using low redundancy of sequencing, typically a single sequencing While single sequencing reactions may have an reaction. accuracy as low as 97%, this nevertheless provides sufficient fidelity for identification of the sequence and design of PCR primers.

Most human genes can be identified by EST sequencing from libraries of cDNA copies of messenger RNAs. However, some genes are expressed only at specific times during embryonic development, or only in small amounts in a few specific cell types. Other genes have mRNAs that are degraded very quickly by the cell in which they are expressed. If any of these are the case, transcripts of the gene will not be represented in cDNA libraries so the gene will not be identifiable by EST sequencing. A new method called "exon amplification", however, can be used to isolate and identify transcripts of such genes.

Exon amplification works by artificially expressing part or all of a gene that is contained in a cloned fragment of genomic DNA such as a cosmid or yeast artificial chromosome (YAC). The gene is cloned into a special vector, designed at MIT, that uses control elements from virus genes to express

5

10

15

20

25

30

35

-11-

the protein-coding exons of the human gene of interest. Exon trapping shows considerable promise as a general technique for identifying those genes in the human genome that cannot found by cDNA cloning and EST sequencing. amplification will also be useful for identifying the genes in regions of genomic DNA to which disease genes have been The exon amplification method can be used directly with the cosmid and YAC clones frown human chromosomes that are being obtained by both NIH and DOE supported human genome ESTs comprise DNA sequences corresponding to a portion of nuclear encoded messenger RNA. An EST is of sufficient length to permit: (1) amplification of the specific sequence from a cDNA library, e.g., by polymerase chain reaction (PCR); (2) use of a synthetic polynucleotide corresponding to a partial or complete sequence of the EST as a hybridization probe of a cDNA library, generally having 30 - 50 base pairs; or (3) unique designation of the pure cDNA clone from which the EST was derived (the EST clone) for use as a hybridization probe of a cDNA library. Preferably, ESTderived primer pairs and sequences amplify or detectably hybridize to a sequence from a genomic library.

It has been found that sufficient information contained in the 150-400 base ESTs from one sequencing run to effect preliminary identification and exact chromosome mapping. Accordingly, the ESTs disclosed herein are generally at least 150 base pairs in length. The length of an EST is determined by the quality of sequencing data and the length of the cloned cDNA. Raw data from the automated sequencers is edited to remove low quality sequence at the end of the sequencing run. High quality sequences (usually a result of sequencing templates without excessive salt contamination) generally give about 400 bp of reliable sequence data; other sequences give fewer bases of reliable data. A 150 bp EST is long enough to be translated into a 50 amino acid peptide sequence. This length is sufficient to observe similarities when they exist in a database search. Furthermore, 150 bp is

10

15

20

35

long enough to design PCR primers from each end of the sequence to amplify the complete EST. Sequences shorter than 150 bp are difficult to purify and use following PCR amplification. Furthermore, a 150 bp polynucleotide is likely to give a very strong signal with low background in a screen of a genomic library.

Finally, it is highly unlikely that a sequence of the same 150 bp exists in any genes in the genome besides the one tagged by the EST. Some closely related gene family members have very similar nucleotide sequences, but no examples of pairs of human genes with long segments of identical sequence have been reported to date. For instance, there are three known β -tubulin genes in humans. Several ESTs were found that matched one or another of these tubulin genes, but several new members of this gene family were also found and could be clearly distinguished from the three known members. ESTs that match perfectly to several different genes can be detected by hybridizing to chromosomes: if many chromosomal loci are observed, the sequence (or a close variant) is present in more than one gene. This problem can be circumvented by using the 3'-untranslated part of the cDNA alone as a probe for the chromosomal location or for the full-length cDNA or gene. The 3'-untranslated region is more likely to be unique within gene families, since there is no evolutionary pressure to conserve a coding function of this region of the mRNA.

As demonstrated in the Examples that follow, ESTs can be used to map the expressed sequence to a particular chromosome. In addition, ESTs can be expanded to provide the full coding regions, as detailed below. In this manner, previously unknown genes can be identified.

While a variety of cDNA libraries can be used to obtain ESTs, human brain cDNA libraries are exemplified and represent a preferred embodiment. Suitable cDNA libraries can be freshly prepared or obtained commercially, e.g., as shown in Examples 1, 2, and 11. The cDNA libraries from the

5

10

15

20

25

30

35

-13-

desired tissue are preferably preprocessed by conventional techniques to reduce repeated sequencing of high and intermediate abundance clones and to maximize the chances of finding rare messages from specific cell populations. Preferably, preprocessing includes the use of defined composition prescreening probes, e.g., cDNA corresponding to mitochondria, abundant sequences, ribosomes, actins, myelin basic polypeptides, or any other known high abundance peptide; these prescreening probes used for preprocessing are generally derived from known ESTs. Other · preprocessing techniques include subtraction. which preferentially reduces the population of certain sequences in the library (e.g., see A. Swaroop et al., Nucl. Acids Res. 19, 1954 (1991)), and normalization, which results in all sequences being represented in approximately proportions in the library (Patanjali et al, Proc. Natl. Acad. Sci. USA 88:1943 (1991)).

The cDNA libraries used in the present method will ideally use directional cloning methods so that either the 5' end of the cDNA (likely to contain coding sequence) or the 3' end (likely to be a non-coding sequence) can be selectively obtained."

Libraries of cDNA can also be generated from recombinant expression of genomic DNA. After they are amplified, ESTs can be obtained and sequenced, e.g., as illustrated in Example 11.

The sequences of the present invention include the specific sequences set forth in the Sequence Listing and designated SEQ ID NO: 1 - SEQ ID NO: 2412. In one aspect of this embodiment, the invention relates to those sequences of SEQ ID NOS: 1 - 2412 that comprise the cDNA coding sequences for polypeptides having less than 95% identity with known amino acid sequences (see Table 2) and more preferably less than 90% or 85% identity. In a second aspect, the invention relates to those sequences of SEQ ID NOS: 1 - 2412 that encode polypeptides having no similarity to known amino acid

-14-

sequences (see Examples that follow). Precisely because they do not contain coding regions and are therefore more unique in their sequence structures, those sequences which meet neither of the preceding criteria can be most useful and are generally preferred for mapping.

Consistent with the NIH mission and its responsibilities to disseminate knowledge and share the tangible fruits of its research, the present inventors have taken a number of steps to facilitate sequence data and clone availability. All EST sequences have been submitted to GenBank (representing an addition equivalent to 7% of the human nucleotides in Release 69 of GenBank, September 1991). The corresponding cDNA clones have been submitted to the American Type Culture Collection and information on clones and sequences has been submitted to the Genome Data Base (Pearson, P. Nucl. Acids Res. 19 (Suppl.): 2237-9 (1991)).

II. Complete Coding Sequences from ESTs

5

10

15

20

25

30

35

The ESTs of the present invention generally represent relatively small coding regions or untranslated regions of human genes. Although most of these sequences do not code for a complete gene product, the ESTs of the present invention are highly specific markers for the corresponding complete coding regions. The ESTs are of sufficient length that they will hybridize, under stringent conditions, only with DNA for that gene to which they correspond. stringent conditions comprise conditions, for example, where at least 95%, preferably at least 97% or 98% identity (base pairing), is required for hybridization. This property permits use of the EST to isolate the entire coding region and even the entire sequence. Therefore, only routine laboratory work is necessary to parlay the unique EST sequence into the corresponding unique complete gene sequence.

Thus, each of the ESTs of the present invention "corresponds" to a particular unique human gene. Knowledge

10

15

20

25

30

of the EST sequence permits routine isolation and sequencing of the complete coding sequence of the corresponding gene. The complete coding sequence is present in a full-length cDNA clone as well as in the gene carried on genomic clones. Therefore, each EST "corresponds" to a cDNA (from which the EST was derived), a complete genomic gene sequence, a polypeptide coding region (which can be obtained either from the cDNA or genomic DNA), and a polypeptide or amino acid sequence encoded by that region.

The first step in determining where an EST is located in the cDNA is to analyze the EST for the presence of coding sequence, e.g., as described in Example 14. The CRM program predicts the extent and orientation of the coding region of a sequence. Based on this information, one can infer the presence of start or stop codons within a sequence and whether the sequence is completely coding or completely non-coding. If start or stop codons are present, then the EST can cover both part of the 5'-untranslated or 3'-untranslated part of the mRNA (respectively) as well as part of the coding sequence. If no coding sequence is present, it is likely that the EST is derived from the 3'-untranslated sequence due to its longer length and the fact that most cDNA library construction methods are biased toward the 3' end of the mRNA.

One general procedure for obtaining complete sequences from ESTs is as follows:

- 1. Purify selected human DNA from an EST clone (the cDNA clone that was sequenced to give the EST), e.g., by endonuclease digestion using ECOR1, gel electrophoresis, and isolation of the aforementioned clone by removal from low-melting agarose gel.
- 2. Radiolabel the isolated insert DNA, e.g., with ³²P labels, preferably by nick translation or random primer labeling.
- 35 3. Use the labeled EST insert as a probe to screen a lambda phage cDNA library or a plasmid cDNA library.

15

20

25.

30

35

- 4. Identify colonies containing clones related to the probe cDNA and purify them by known purification methods.
- 5. Nucleotide sequence the ends of the newly purified clones to identify full length sequences.
- 6. Perform complete sequencing of full length clones by Exonuclease III digestion or primer walking. Northern blots of the mRNA from various tissues using at least part of the EST clone as a probe can optionally be performed to check the size of the mRNA against that of the purported full length cDNA.

An EST is a specific tag for a messenger RNA molecule. The complete sequence of that messenger RNA, in the form of cDNA, can be determined using the EST as a probe to identify a cDNA clone corresponding to a full-length transcript, followed by sequencing of that clone. The EST or the full-length cDNA clone can also be used as a probe to identify a genomic clone or clones that contain the complete gene including regulatory and promoter regions, exons, and introns.

ESTs are used as probes to identify the cDNA clones from which an EST was derived. ESTs, or portions thereof, can be nick-translated or end-labelled with P32 using polynucleotide kinase using labelling methods known to those with skill in the art (Basic Methods in Molecular Biology, L.G. Davis, M.D. Dibner, and J.F. Battey, ed., Elsevier Press, NY, 1986). lambda library can be directly screened with the labelled ESTs of interest or the library can be converted en masse to pBluescript (Stratagene, La Jolla, California) to facilitate bacterial colony screening. Both methods are well known in the art. Briefly, filters with bacterial colonies containing the library in pBluescript or bacterial lawns containing lambda plaques are denatured and the DNA is fixed to the filters. The filters are hybridized with the labelled probe using hybridization conditions described by Davis et al. The ESTs, cloned into lambda or pBluescript, can be used as positive controls to assess background binding and to adjust

the hybridization and washing stringencies necessary for accurate clone identification. The resulting autoradiograms are compared to duplicate plates of colonies or plaques; each exposed spot corresponds to a positive colony or plaque. The colonies or plaques are selected, expanded and the DNA is isolated from the colonies for further analysis and sequencing.

5

10

.15

20

25

30

35

The ESTs can additionally be used to screen Northern blots of mRNA obtained from various tissues or cell cultures, including the tissue of origin of the EST clone. Northern analysis will most often produce one to several positive bands. The bands can be selected for further study based on the predicted size of the mRNA.

Positive cDNA clones in phage lambda are analyzed to determine the amount of additional sequence they contain using PCR with one primer from the EST and the other primer from the vector. Clones with a larger vector-insert PCR product than the original EST clone are analyzed by restriction digestion and DNA sequencing to determine whether they contain an insert of the same size or similar as the mRNA size on a Northern blot.

Once one or more overlapping cDNA clones are identified, the complete sequence of the clones can be determined. The preferred method is to use exonuclease III digestion (McCombie, W.R, Kirkness, E., Fleming, J.T., Kerlavage, A.R., Iovannisci, D.M., and Martin-Gallardo, R., Methods: 3: 33-40, 1991). A series of deletion clones is generated, each of which is sequenced. The resulting overlapping sequences are assembled into a single contiguous sequence of high redundancy (usually three to five overlapping sequences at each nucleotide position), resulting in a highly accurate final sequence.

A similar screening and clone selection approach can be applied to obtaining cosmid or lambda clones from a genomic DNA library that contains the complete gene from which the EST was derived (Kirkness, E.F., Kusiak, J.W., Menninger, J.,

15

20

25

30

35

Gocayne, J.D., Ward, D.C., and Venter, J.C. Genomics 10: 985-995 (1991). Although the process is much more laborious, these genomic clones can be sequenced in their entirety also. A shotgun approach is preferred to sequencing clones with inserts longer than 10 kb (genomic cosmid and lambda clones). In shotgun sequencing, the clone is randomly broken into many small pieces, each of which is partially sequenced. The sequence fragments are then aligned to produce the final contiguous sequence with high redundancy. An intermediate approach is to sequence just the promoter region and the intron-exon boundaries and to estimate the size of the introns by restriction endonuclease digestion (ibid.).

Using the sequence information provided herein, the polynucleotides of the present invention can be derived from natural sources or synthesized using known methods. sequences falling within the scope of the present invention are not limited to the specific sequences described, but include human allelic and species variations thereof and portions thereof of at least 15-18 bases. (Sequences of at least 15-18 bases can be used, for example, as PCR primers or as DNA probes.) In addition, the invention includes the coding sequence associated with the entire polynucleotide sequence of bases described in the Sequence Listing, as well as portions of the entire coding sequence of at least 15-18 bases and allelic and species variations thereof. Furthermore, to accommodate codon variability, the invention includes sequences coding for the same amino acid sequences as do the specific sequences disclosed herein. Finally, although the error rate in the automated sequencing used in the present invention is small, there remains some chance of error. Therefore, claims to particular sequences should not be so narrowly construed as to require inclusion of erroneously identified bases or to exclude corrections.

Any specific sequence disclosed herein can be readily screened for errors by resequencing each EST in both directions (i.e., sequence both strands of cDNA).

The sequences, constructs, vectors, clones, and other materials comprising the present invention can advantageously be in enriched or isolated form. As used herein, "enriched" means that the concentration of the material is at least about 2, 5, 10, 100, or 1000 times its natural concentration (for example), advantageously 0.01%, by weight, preferably at least about 0.1% by weight. Enriched preparations of about 0.5%, 1%, 5%, 10%, and 20% by weight are also contemplated. Further, removal of clones corresponding to ribosomal RNA and "housekeeping" genes and clones without human cDNA inserts results in a library that is "enriched" in the desired clones.

5

10

15

20 .

25

30

35

The term "isolated" requires that the material be removed from its original environment (e.g., the natural environment if it is naturally occurring). For example, a naturally-occurring polynucleotide present in a living animal is not isolated, but the same polynucleotide, separated from some or all of the coexisting materials in the natural system, is isolated.

It is also advantageous that the sequences be in purified form. The term "purified" does not require absolute purity; rather, it is intended as a relative definition. Individual EST clones isolated from a cDNA library have been conventionally purified to electrophoretic homogeneity. The sequences obtained from these clones could not be obtained directly either from the library or from total human DNA. The cDNA clones are not naturally occurring as such, but rather are obtained via manipulation of a partially purified naturally occurring substance (messenger RNA). conversion of mRNA into a cDNA library involves the creation of a synthetic substance (cDNA) and pure individual cDNA clones can be isolated from the synthetic library by clonal selection. Thus, creating a cDNA library from messenger RNA and subsequently isolating individual clones from that library results in an approximately 106-fold purification of the native message. Purification of starting material or

10

15

20

25

30

natural material to at least one order of magnitude, preferably two or three orders, and more preferably four or five orders of magnitude is expressly contemplated.

In a cDNA library there are many species of mRNA represented. Each cDNA clone can be interesting in its own right, but must be isolated from the library before further experimentation can be completed. In order to sequence any specific cDNA, it must be removed and separated (i.e. isolated and purified) from all the other sequences. This can be accomplished by many techniques known to those of skill in the art. These procedures normally involve identification of a bacterial colony containing the cDNA of interest and further amplification of that bacteria. Once a cDNA is separated from the mixed clone library, it can be used as a template for further procedures such as nucleotide sequencing.

Although claims to large numbers of ESTs and corresponding sequences are presented herein, the invention is not limited to these particular groupings of sequences. Thus, individual sequences are considered as applicants' discoveries or inventions, as are subgroupings of sequences. All of the functional subgroupings set forth in the tables define groupings for which separate claims are contemplated as being within the scope of this invention. addition to claims to individual clones, it is intended that the present disclosure also support claims to numerical subgroupings. Thus, subgroupings of 50 ESTs (and corresponding sequences) are contemplated (e.g., SEQ ID NOS 1-50, 51-100, 101-150, etc.) as being within the scope of this invention, as are subgroupings of 5, 10, 25, 100, 200, and 500 ESTs and corresponding sequences.

III. DNA Constructs

The present invention also includes recombinant constructs comprising one or more of the sequences as broadly described above. The constructs comprise a vector, such as

10

15

20

25

30

35

a plasmid or viral vector, into which a sequence of the invention has been inserted, in a sense or antisense orientation. In a preferred aspect of this embodiment, the construct further comprises regulatory sequences, including for example, a promoter, operably linked to the sequence. Large numbers of suitable vectors and promoters are known to those of skill in the art, and are commercially available. The following vectors are provided by way of example. Bacterial: pBs, phagescript, ϕ X174, pBluescript SK, pBs KS, pNH8a, pNH16a, pNH18a, pNH46a (Stratagene); pTrc99A, pKK223-3, pKK233-3, pDR540, pRIT5 (Pharmacia).

Eukaryotic: pWLneo, pSV2cat, pOG44, pXT1, pSG (Stratagene);
pSVK3, pBPV, pMSG, pSVL (Pharmacia).

Promoter regions can be selected from any desired gene using CAT (chloramphenicol transferase) vectors or other vectors with selectable markers. Two appropriate vectors are pKK232-8 and pCM7. Particular named bacterial promoters include lacI, lacZ, T3, T7, gpt, lambda P_R , and trc. Eukaryotic promoters include CMV immediate early, HSV thymidine kinase, early and late SV40, LTRs from retrovirus, and mouse metallothionein-I. Selection of the appropriate vector and promoter is well within the level of ordinary skill in the art.

In a further embodiment, the present invention relates to host cells containing the above-described construct. The host cell can be a higher eukaryotic cell, such as a mammalian cell, or a lower eukaryotic cell, such as a yeast cell, or the host cell can be a procaryotic cell, such as a bacterial cell. Introduction of the construct into the host cell can be effected by calcium phosphate transfection, DEAE dextran mediated transfection, or electroporation (Davis, L., Dibner, M., Battey, I., Basic Methods in Molecular Biology, (1986)).

The constructs in host cells can be used in a conventional manner to produce the gene product coded by the recombinant sequence. Alternatively, the encoded polypeptide

15

20

25

30

35

can be synthetically produced by conventional peptide synthesizers.

Certain ESTs have already been preliminarily categorized by analogy to related sequences in other organisms (see Table 2). Table 10 of Example 10 categorizes particular ESTs broadly as metabolic, regulatory, and structural sequences where known. Constructs comprising genes or coding sequences corresponding to each of these categories are, therefore, specifically and individually contemplated.

Table 11 more particularly separates 127 new ESTs into 13 categories using a different criteria. These are genes related to cell surface; developmental control; energy and phosphatase; oncogenes; metabolism; kinase metabolism-related polypeptides; peptidases and peptidase inhibitors; receptors; structural and cytoskeletal; signal transduction; transporters; transcription, translation, and subcellular localization; and transcription factors. Table 11 further identifies the EST by the particular gene product for which it apparently codes. Each of these categories individually comprises a preferred category of EST, preferred constructs and resulting polypeptide can be prepared from those ESTs or the corresponding complete gene sequence.

IV. ESTs and Corresponding Sequences as Reagents

Each of the cDNA sequences identified herein (and the corresponding complete gene sequences) can be used in numerous ways as polynucleotide reagents. The sequences can be used as diagnostic probes for the presence of a specific mRNA in a particular cell type. In addition, these sequences can be used as diagnostic probes suitable for use in genetic linkage analysis (polymorphisms). Further, the sequences can be used as probes for locating gene regions associated with genetic disease, as explained in more detail below.

The EST and complete gene sequences of the present invention are also valuable for chromosome identification.

-23-

Each sequence is specifically targeted to and can hybridize with a particular location on an individual human chromosome. Moreover, there is a current need for identifying particular sites on the chromosome. Few chromosome marking reagents based on actual sequence data (repeat polymorphisms) presently available for marking chromosomal location. The present invention constitutes a major expansion of available chromosome markers. One hundred ESTS have already been mapped to chromosomes. Using the techniques described in Example 5 or 6, the remaining ESTs and the corresponding complete sequences can similarly be mapped to chromosomes. The mapping of ESTs and cDNAs to chromosomes according to the present invention is an important first step in correlating those sequences with genes associated with disease.

5

10

15

20

25

30

35

Briefly, sequences can be mapped to chromosomes by preparing PCR primers (preferably 15-25 bp) from the ESTs. Computer analysis of the ESTs is used to rapidly select primers that do not span more than one exon in the genomic DNA, thus complicating the amplification process. These primers are then used for PCR screening of somatic cell hybrids containing individual human chromosomes. Only those hybrids containing the human gene corresponding to the EST will yield an amplified fragment.

PCR mapping of somatic cell hybrids is a rapid procedure for assigning a particular EST to a particular chromosome. Three or more clones can be assigned per day using a single thermal cycler. Using the present invention with the same oligonucleotide primers, sublocalization can be achieved with panels of fragments from specific chromosomes or pools of large genomic clones in an analogous manner. Other mapping strategies that can similarly be used to map an EST to its chromosome include in situ hybridization, prescreening with labeled flow-sorted chromosomes and preselection hybridization to construct chromosome specific CDNA libraries. Results of mapping ESTs to chromosomal segments are listed in Tables 3 and 4.

10

20

25

30

35

Fluorescence in situ hybridization (FISH) of a cDNA clone to a metaphase chromosomal spread can be used to provide a precise chromosomal location in one step. This technique can be used with cDNA as short as 500 or 600 bases; however, clones larger than 2,000 bp have a higher likelihood of binding to a unique chromosomal location with sufficient signal intensity for simple detection. FISH requires use of the clone from which the EST was derived, and the longer the better. 2,000 bp is good, 4,000 is better, and more than 4,000 is probably not necessary to get good results a reasonable percentage of the time. For a review of this technique, see Verma et al., Human Chromosomes: a Manual of Basic Techniques. Pergamon Press, New York (1988).

Reagents for chromosome mapping can be used individually (to mark a single chromosome or a single site on that chromosome) or as panels of reagents (for marking multiple sites and/or multiple chromosomes). Reagents corresponding to noncoding regions of the genes actually are preferred for mapping purposes. Coding sequences are more likely to be conserved within gene families, thus increasing the chance of cross hybridizations during chromosomal mapping (see Tables 8 and 9).

Once a sequence has been mapped to a precise chromosomal location, the physical position of the sequence on the chromosome can be correlated with genetic map data. (Such data are found, for example, in V. McKusick, Mendelian Inheritance in Man (available on line through Johns Hopkins University Welch Medical Library).) The relationship between genes and diseases that have been mapped to the same chromosomal region are then identified through linkage analysis (coinheritance of physically adjacent genes).

Next, it is necessary to determine the differences in the cDNA or genomic sequence between affected and unaffected individuals. If a mutation is observed in some or all of the affected individuals but not in any normal individuals, then

the mutation is likely to be the causative agent of the disease.

With current resolution of physical mapping and genetic mapping techniques, a cDNA precisely localized to a chromosomal region associated with the disease could be one of between 50 and 500 potential causative genes. (This assumes 1 megabase mapping resolution and one gene per 20 kb.)

5

10

15

20

25

30

35

Comparison of affected and unaffected individuals generally involves first looking for structural alterations in the chromosomes, such as deletions or translocations that are visible from chromosome spreads or detectable using PCR based on that cDNA sequence. Ultimately, complete sequencing of genes from several individuals is required to confirm the presence of a mutation and to distinguish mutations from polymorphisms.

In addition to the foregoing, the sequences of the invention, as broadly described, can be used to control gene expression through triple helix formation or antisense DNA or RNA, both of which methods are based on binding of a polynucleotide sequence to DNA or RNA. Polynucleotides suitable for use in these methods are usually 20 to 40 bases in length and are designed to be complementary to a region of the gene involved in transcription (triple helix - see Lee et al, Nucl. Acids Res. 6: 3073 (1979); Cooney et al, Science 241: 456 (1988); and Dervan et al, Science 251: 1360 (1991)) or to the mRNA itself (antisense - Okano, J. Neurochem. 56: 560 (1991); Oligodeoxynucleotides as Antisense Inhibitors of Gene Expression, CRC Press, Boca Raton, FL (1988)). helix formation optimally results in a shut-off of RNA transcription from DNA, while antisense RNA hybridization blocks translation of an mRNA molecule into polypeptide. Both techniques have been demonstrated to be efficient in model systems. Information contained in the sequences of the present invention is necessary for the design of an antisense or triple helix oligonucleotide.

15

20

25

30

35

The present invention is also useful tool in gene therapy, which requires isolation of the disease-associated gene in question as a prerequisite to the insertion of a normal gene into an organism to correct a genetic defect. He high specificity of the cDNA probes according to this invention have promise of targeting such gene locations in a highly accurate manner.

The sequences of the present invention, as broadly defined, are also useful for identification of individuals from minute biological samples. The United States military, for example, is considering the use of restriction fragment length polymorphism (RFLP) for identification of its personnel. In this technique, an individual's genomic DNA is digested with one or more restriction enzymes, and probed on a Southern blot to yield unique bands for identifying personnel. This method does not suffer from the current limitations of "Dog Tags" which can be lost, switched, or stolen, making positive identification difficult. sequences of the present invention are useful as additional DNA markers for RFLP.

However, RFLP is a pattern based technique, which does not directly focus on the actual DNA sequence of the individual. The sequences of the present invention can be used to provide an alternative technique that determines the actual base-by-base DNA sequence of selected portions of an individual's genome. These sequences can be used to prepare PCR primers for amplifying and isolating such selected DNA. One can, for example, take an EST of the invention and prepare two PCR primers from the 5' and 3' ends of the EST. These are used to amplify an individual's DNA, corresponding to the EST. The amplified DNA is sequenced.

Panels of corresponding DNA sequences from individuals, made this way, can provide unique individual identifications, as each individual will have a unique set of such DNA sequences, due to allelic differences. The sequences of the present invention can be used to particular advantage to

10

15

20

25

30

35

obtain such identification sequences from individuals and from tissue, as explained in Examples 12 - 14.

The EST sequences from Examples 1 and 2 and the complete sequences from Example 13 uniquely represent portions of the human genome. Allelic variation occurs to some degree in the coding regions of these sequences, and to a greater degree in the noncoding regions. It is estimated that allelic variation between individual humans occurs with a frequency of about once per each 500 bases. Each of the ESTs or complete coding sequences comprising a part of the present invention can, to some degree, be used as a standard against DNA which from an individual can be compared identification purposes. Because greater numbers polymorphisms occur in the noncoding regions, fewer sequences are necessary to differentiate individuals. The noncoding sequences of Table 9 for example, could comfortably provide positive individual identification with a panel of perhaps 100 to 1,000 primers which each yield a noncoding amplified sequence of 100 bp. If predicted coding sequences, such as those from Table 6, are used, a more appropriate number of primers for positive individual identification would be 500-2,000.

If a panel of reagents from ESTs or complete sequences of this invention is used to generate a unique ID database for an individual, those same reagents can later be used to identify tissue from that individual. Positive identification of that individual, living or dead can be made from extremely small tissue samples.

Another use for DNA-based identification techniques is in forensic biology. PCR technology can be used to amplify DNA sequences taken from very small biological samples such as tissues, e.g., hair or skin, or body fluids, e.g., blood, saliva, semen, etc. In one prior art technique, gene sequences are amplified at specific loci known to contain a large number of allelic variations, for example the DQ α class II HLA gene (Erlich, H., PCR Technology, Freeman and Co.

10

15

20

25

30

(1992)). Once this specific area of the genome is amplified, it is digested with one or more restriction enzymes to yield an identifying set of bands on a Southern blot probed with DNA corresponding to the DQ α class II HLA gene.

The sequences of the present invention can be used to provide polynucleotide reagents specifically targeted to additional loci in the human genome, and can enhance the reliability of DNA-based forensic identifications. sequences targeted to noncoding regions (see, e.g., Tables 8 and 9) are particularly appropriate. As mentioned above, actual base sequence information can be used for identification as an accurate alternative to patterns formed by restriction enzyme generated fragments. Reagents for obtaining such sequence information are within the scope of the present invention. Such reagents can comprise complete ESTs or corresponding coding regions, or fragments of either of at least 15 bp, preferably at least 18 bp.

There is also a need for reagents capable of identifying the source of a particular tissue. Such need arises, for example, in forensics when presented with tissue of unknown origin. Appropriate reagents can comprise, for example, DNA probes or primers specific to particular tissue prepared from the ESTs or complete sequences of the present invention. Panels of such reagents can identify tissue by species and/or by organ type. In a similar fashion, these reagents can be used to screen tissue culture for contamination.

V. Production of Polypeptide Corresponding to ESTs

As previously explained, each EST corresponds not only to a coding region, but also to a polypeptide. Once the coding sequence is known, or the gene is cloned which encodes the polypeptide, conventional techniques in molecular biology can be used to obtain the polypeptide.

At the simplest level, the amino acid sequence encoded by the polynucleotide sequence can be synthesized using commercially available peptide synthesizers. This is

10

15

20

25

30

particularly useful in producing small peptides and fragments of larger polypeptides. (Fragments are useful, for example, in generating antibodies against the native polypeptide.)

Alternatively, the DNA encoding the desired polypeptide can be inserted into a host organism and expressed. organism can be a bacterium, yeast, cell line. or multicellular plant or animal. The literature is replete with examples of suitable host organisms and expression techniques. For example, naked polynucleotide (DNA or mRNA) can be injected directly into muscle tissue of mammals, where it is expressed. This methodology can be used to deliver the polypeptide to the animal, or to generate an immune response against a foreign polypeptide. Wolff, et al., 247:1465 (1990); Felgner, et al., Nature 349:351 (1991). Alternatively, the coding sequence, together with appropriate regulatory regions (i.e., a construct), can be inserted into a vector, which is then used to transfect a cell. The cell (which may or may not be part of a larger organism) then expresses the polypeptide. (See Example 25.)

Antibodies generated against the polypeptide corresponding to a sequence of the present invention can be obtained by direct injection of the naked polypeptide into an animal (as above) or by administering the polypeptide to an animal, preferably a nonhuman. The antibody so obtained will then bind the polypeptide itself. In this manner, even a sequence encoding only a fragment of the polypeptide can be used to generate antibodies binding the whole native polypeptide. Such antibodies can then be used to isolate the polypeptide from tissue expressing that polypeptide. Moreover, a panel of such antibodies, specific to a large number of polypeptides, can be used to identify and differentiate such tissue.

VI. Examples

Certain aspects of the present invention are described in greater detail in the non-limiting Examples that follow.

EXAMPLE 1

<u>Clone Selection: First set</u>

5

10

15

20

25

30

METHODOLOGY:

With reference to the data presented in Table 1, lambda libraries were converted en masse to pBluescript plasmids, transfected into E. coli XL1-Blue cells, and plated on X-gal/IPTG/ampicillin plates. A total of 1058 clones were picked at random from three human brain cDNA libraries: fetal brain, two-year-old hippocampus, and two-year-old temporal cortex (Stratagene catalog #936206, 936205, 935, respectively. Stratagene, 11099 N. Torrey Pines Rd., La Jolla, CA 92037). An analysis of these clones is summarized in Table I (see below) In addition, clones selected from the hippocampus library were also analyzed after subtractive hybridization with the fibroblast library. These results are listed in the "Hippocampus Subtracted" column of Table 1. Templates for DNA sequencing were PCR products or plasmids prepared by the alkaline lysis method. About half of the templates prepared by PCR failed to yield an amplified fragment suitable for sequencing. This was primarily due to use of PCR conditions that minimized the need for further purification of the product but also selected against amplification of long inserts (5 μ l fresh or frozen overnight culture of E. coli carrying the pBluescript plasmid, 7.5 μM each dNTP, and 0.1 μM each primer for 35 cycles: 40 sec; 55°C, 40 sec; 72°C, 90 sec). A further percentage of the PCR-generated templates failed to sequence, largely due to primer-dimer or other amplification artifacts. Qiagen™ columns improved the percentage of plasmid templates, increasing the yields of usable sequence from about 60% with a standard alkaline lysis protocol to over 90%. Overall, 117 PCR-generated templates and 497 plasmid templates resulted in usable sequence. Dideoxy chain termination sequencing reactions were performed with fluorescent dye-labeled M13

35

universal or reverse primers. After a cycle sequencing protocol, carried out in a Perkin-Elmer thermal cycler, sequencing reactions were run on an Applied Biosystems, Inc. (Foster City, CA) 373A automated DNA sequencer. (Cycle sequencing was performed in a Perkin Elmer Thermal Cycler for 15 cycles of 95°C, 30 sec; 60°C, 1 sec; 70°C, 60 sec and 15 cycles of 95°C, 30 sec; 70°C, 60 sec with the Applied Biosystems, Inc. Taq Dye Primer Cycle Sequencing Core Kit protocol). Some sequencing reactions were performed on an ABI robotic workstation (Cathcart, Nature 347: 310 (1990) hereby incorporated by reference).

RESULTS:

5

10

15

20

25

30

35

Singe-run DNA sequence data were obtained from 609 randomly chosen cDNA clones. The number of clones sequenced from each library is summarized in Table 1. Double-stranded cDNA clones in the pBluescript vector were sequenced by a cycle sequencing protocol with dye-labeled primers and Applied Biosystems, Inc. 373A DNA Sequences. The average length of usable sequence was 397 bases with a standard deviation of 99 bases.

Subtractive hybridization has been used successfully to reduce the population of highly represented sequences in a cDNA library by selectively removing sequences shared by another library. (Schmid and Girou, Neurochem. 48: 307 (1987); Fargnoli et al, Anal. Biochem. 187: 364 (1990); Duguid and Dinauer, Nucl. Acids. Res. 18: 2789 Schweinfest, et al, Genet. Anal. Techn. Appl. 7: 64 (1990); Travis and Sutcliffe, Proc. Natl. Acad. Sci. USA 85: 1696 (1988); Kato, Eur. J. Neurosci. 2: 704 (1990)). Subtractive hybridization was therefore tested as a way of enhancing the number of brain-specific clones in the hippocampus library by hybridizing the hippocampus library with a WI38 human lung fibroblast cell line cDNA library and removing the common sequences (Schweinfest et al, G net. Anal. Techn. Appl. 7: 64 (1990); Sive and St. John, Nucl. Acids Res. 16: 10937

(1988)). Clones from this subtraction are listed in the column "Hippocampus Subtracted" in Table 1.

The EST sequences from this Example 1 are identified as SEQ ID NOs 1-315.

-33-

TABLE 1. cDNA Library Composition Determined By Random Clone Sequencing

	int		'n		60		0		_	س	
Cortex	Percent		7.	0	13.	0	, ry	. ~	33.	32.5	
Tempora	Number		•	0	=	0	•	• • •	27	5 2	
Fetal Brain	Percent		7.9	15.8	0	10.5	13.2	52.6	0	0	
	Number		.	•	0	7	'n	2	0	0 .	
Hippocampus Subtracted	Percent	,	9.8	12.2	6.0	6.0	6.0	37.9	20.7	5.6	
Hippocampu	Number	;	9	*		7	~	7,	*	m _.	
H i ppocampus	Percent		12.8	10.4	2.7	8.6	8.6	42.8	14.1	0.3	
H ppo	Number	:	84	39	2	35	35	160	53	-	
		uman	nes	ine-1, etc.		res	ter				
	EST Category	Databases MatchHuman	Mitochondrial Ger	Repeats: Alu, Li	Ribosomal RNA	Other Nuclear Ger	Database MatchOth	No Database Match	poly A Insert	No Insert	

10

15

20

25

30

EXAMPLE 2

Sequencing of Additional ESTs: Second set

Over 2600 additional cDNA clones have been isolated, partially sequenced and screened. The clones were isolated from four human brain cDNA libraries. The new sequences thus discovered, together with the 315 brain ESTs from Example 1, correspond to over 2400 new human genes. These data represent an approximate doubling of the number of human genes identified by DNA sequencing.

Specifically, four cDNA libraries were used as sources of clones for sequencing. Human hippocampus and fetal brain preparation, libraries, plasmid template sequencing reactions, and automated sequencing were performed as described (Adams, M.D., Kelley, J.M., Gocayne, J.D., Dubnick, M., Polymeropoulos, M.H., Xiao, H., Merril, C.R., Wu, A., Olde, B., Moreno, R.F., Kerlavage, A.R., McCombie, W.R., & Venter, J.C. Science, 252: 1651-56 (1991)). A pooled probe consisting of inserts from 10 different EST clones with sequences that matched either mitochondrial genes or the 185 or 28S ribosomal RNAs was used to prescreen a gridded filter array of the hippocampus library; nonhybridizing clones are referred to as the "prescreened library". Another fetal. brain library was constructed by and was a gift from Bento Soares (Columbia University). A directionally-cloned library was prepared using the method of Rubenstein, (Rubenstein, J., Elizabeth, A., Brice, A., Ciaranello, R., Denney, D., Porteus, M. & Usdin, T. Nucl. Acids Res. 18: 4833-4842) using human adult brain mRNA purchased from Clontech (Palo Alto, CA; Catalogue # 6516-1). Of 482 clones analyzed by restriction enzyme digestion, 33% contained inserts at least 1500 base pairs in length. hippocampus and fetal brain library totals include data from Adams et al Science 252: 1651.

Sequences of nuclear-encoded cDNAs that did not include interspersed repeats (Schmid, C. W. & Jelinek, W. R. Science

10

15

20

25

30

216: 1065-1070 (1982); Paulson, K. E., Deka, N., Schmid, C. W., Misra, R., Schlinder, C. W., Rush, M. G., Kadyk, L., & Leinwand, L. Nature 316: 359-361 (1985); Fanning, T. G. & Biochem. Biophys. Acta 910: 203-212 (1987)) Singer, M. F. were searched against all of GenBank and, in 6-frame translation, against a comprehensive, non-redundant peptide database using the network BLAST (Altschul, S. F., Gish, W., Miller, W., Myers, E.W., & Lipman, D. J. Mol. Biol. 215: (1990)) server at the National Center Biotechnology Information. BLAST output was parsed, and an interactive alignment editor was used to select which matches, if any, from each search to record in a relational EST database, which was developed to track sequencing, identification, tissue localization, physical mapping, and the public distribution of the clones, mapping and sequence data. For significant similarities, a putative gene name and Protein Identification Resource (PIR) gene · family identification (Barker, W., George, D., Hunt, L., Garavelli, J. Nucl. Acids Res. 19 (Suppl): 2231-2236 (1991)) for the EST were assigned. ESTs without significant matches using BLAST were searched in translation against PIR using FASTA. Ten additional marginal matches were found. A total of 2300 new EST sequences comprising 765,505 nucleotides from the current data set have been submitted to GenBank and assigned accession numbers M77851-M79278 and M85308-M86179. All ESTs except those multiply representing actin, tubulin, and myelin basic protein clones were submitted. accession numbers of cDNA clones from which ESTs were derived are 77501-78999 and 81000-81756. The Genome Data Base expressed D-segment numbers for these clones are DOS1E -D0S2422E. The ESTs from this Example are identified herein as SEQ ID NOs 316-2407.

10

15

20

.25

30

35

EXAMPLE 3

EST Characterization: First Set

ESTs including SEQ ID NOs 1-315 were analyzed as follows. Initially, the EST sequences were examined for similarities in the GenBank nucleic acid database (GenBank Release 65.0), Protein Information Resource Release 26.0 (PIR), and ProSite (MacPattern from the EMBL data library, Fuchs R. Comput. Appl. Biosci. 7: 105 (1990) Release 5.0 were used). BLAST was used to search Genbank and the PIR (both maintained by the National Center for Biotechnology Information) ESTs without exact GenBank matches translated in all six reading frames and each translation was compared with the protein sequence database PIR and the ProSite protein motif database. Comparisons with the ProSite motif database were done by means of the program MacPattern from the EMBL Data Library. GenBank and PIR searches were conducted with the "basic local alignment search tool" programs for nucleotide (BLASTN) and peptide comparisons (Altschul et al, J. Mol. Biol. 215: 403 (1990)). searches were run on the National Center Biotechnology Information BLAST network service. The BLAST programs contain a very rapid database-searching algorithm that searches for local areas of similarity between two sequences and then extends the alignments on the basis of defined match and mismatch criteria. The algorithm does not consider the potential gaps to improve the alignment, thus sacrificing some sensitivity for a 6-80 fold increase in speed over other database-searching programs such as FASTA (Pegarson and Lipman, Proc. Natl. Acad. Sci. USA, 85: 2444 (1988)).

Sequence similarities identified by the BLAST programs were considered statistically significant with a Poisson P-value than 0.01. The Poisson P-value less than the probability of as high a score occurring by chance given the number of residues in the query sequence and the database.

WO 93/16178 PCT/US93/01294⁻

After the BLASTN search, 30 unmatched ESTs were compared against GenBank by FASTA to determine if significant matches were missed due to the use of BLASTN for the database search. No additional statistically significant matches were found. Statistical significance does not necessarily mean functional similarity; some of the reported matches may indicate the presence of a conserved domain or motif or simply a common protein structure pattern. Those ESTs identified as fully corresponding to known human genes or proteins are not included in this disclosure. Statistically significant matches are reported in Table 2, together with the length and percent identity or similarity of each alignment.

5

10

15

20

25

30

35

On the basis of database searches, 609 EST sequences were classified into eight groups as shown in Table 1 (see Example 1 above). Four groups, with 197 or 32% of the sequences, consist of matches to human sequences: repetitive elements, mitochondrial genes, ribosomal RNA genes, and other nuclear genes. Forty-eight (8%) of the sequences matched non-human entries in GenBank or PIR while 230 (38%) had no significant matches. The remaining 134 (22%) sequences contained no insert or consisted entirely of polyA between the EcoRI cloning sites.

Thirty-six ESTs matched previously sequenced human nuclear genes with more than 97% identity. Four of these ESTs are from genes encoding enzymes involved in maintaining metabolic energy, including ADP/ATP translocase, aldolase C, hexokinase, and phosphoglycerate kinase. Human homologs of genes for the bovine mitochondrial ATP synthase $F_0 \mbox{$\mathbb{B}$-subunit}$ and porcine aconitase were also found (Table 2). specific cDNAs included synaptophysin, glial fibrillary acidic protein (GFAP), and neurofilament light chain. least six ESTs are from genes encoding proteins involved in signal transduction: 2',3'-cyclic nucleotide 3 ' phosphodiesterase (2 ESTs), calmodulin, c-erbs- α -2, $G_S\alpha$, and Na^+/K^+ ATPase α -subunit. Other ESTs were matches to genes for ubiquitous structural proteins -- actins, tubulins, and

fodrin (non-erythroid spectrin). ESTs also document the presence in the hippocampus cDNA library of the ret proto-oncogene, the ras-related gene rhoB, and one of the chromosome 22 breakpoint cluster region transcripts. Eight ESTs are from genes known to be associated with genetic disorders (Online Mendelian Inheritance in Man). More than half of the human-matched ESTs from Example 1 have been mapped to chromosomes, indicating the bias of GenBank entries toward well-studied genes and proteins.

10

5

ESTs without significant GenBank matches were also compared to the ProSite database of recognized protein motifs. Not counting post-translational-modification signatures, fifty-four sequences contained motifs from the database. Some patterns, particularly the "leucine zipper", are found in scores or hundreds of proteins that do not share the functional property implied by the presence of the motif.

15

20

25

Similarities to sequences from other organisms were also detected in the BLAST searches of GenBank and PIR (Table 2). Several ESTs displayed similarity to "housekeeping" genes, including the ribosomal proteins S10 and L30 (rat) and the above glycolytic enzymes. EST00257 (SEQ ID NO:77) shows strong nucleotide sequence similarity to the squid (67%) and Drosophila (70.4%) kinesin heavy chain. Kinesin was first described as a microtubule-associated motor protein involved in organelle transport in the squid giant axon (Vale et al, Cell 42: 39 (1985)). Six oncogene-related sequences were also among the cDNA clones sequenced. EST00299 (SEQ ID NO:180) and EST00283 (SEQ ID NO:271) show similarity to several ras-related genes and EST00248 (SEQ ID NO:102) matched the 3' untranslated region of the bovine substrate of botulinum toxin ADP-ribosyltransferase. Similarities with an S. cerevisiae RNA polymerase subunit and Torpedo electromotor neuron-associated protein were also observed. Two ESTs may represent new members of known human gene families: EST00270 matched the three ß-tubulin genes with 88-91% identity and

30

35

10

15

20

25

30

35

EST00271 (SEQ ID NO:248) matched α -actinin with 85% identity at the nucleotide level.

Among the most interesting of the primary sequence relationships was the similarity of ESTs to the Drosophila genes Notch and Enhancer of split. Nucleotide and peptide alignments of EST00256 (SEQ ID NO:188) and EST00259 (SEQ ID NO:227) with the Drosophila genes have been demonstrated. Both genes are part of a signal cascade encoded by the "neurogenic" genes that are involved in the differentiation of neuronal and epidermal cell lineages in the neuroectoderm of the developing Drosophila embryo (Campos-Ortega, Trends in Neuro. Sci. 11: 400 (1988)). It has been proposed that the Enhancer of split protein interacts with a membrane protein is the product of the Notch gene to convert signal into an altered pattern of gene developmental expression (id. J. Mol. Biol. 215: 403 (1990)). (SEQ ID NO:188) matches near the 5' end of the Enhancer of split coding sequence, away from the mammalian G protein & subunit- and yeast cdc4-like elements (Hartley et al, Cell 55: 785 (1988); Klambt et al. EMBO J. 8: 203 (1989)). Part of the EST00259 (SEQ ID NO:227) match to Notch in the cdc10/SW16 region that is similar to three cell-cycle control genes in yeast and is tightly conserved in the Xenopus Notch homolog, Xotch. In Drosophila, Enhancer of split is absolutely required for formation of epidermal tissue. Notch contains several epidermal growth factor-like repeats and appears to play a general role in cell-cell communication during development (Banerjee and Zipursky, Neuron 4:177 (1990)).

Seven genes were represented by more than one EST. Comparisons of all the ESTs against one another revealed two overlaps of unknown ESTs: EST00233 (SEQ ID NO:32) and EST00234 (SEQ ID NO:8) match in opposite orientations and EST00235 (SEQ ID NO:204) and EST00236 (SEQ ID NO:148) match in the same orientation beginning at the same nucleotide. Five human genes were represented by more than one EST: ß-

10

15

20

25

30

actin (3), λ -actin (2), α -tubulin (2), α -2-macroglobulin (2), and 2'3'-cyclic-nucleotide-3'-phosphodiesterase (2). Those few instances where two or more ESTs represent different portions of a single cDNA can be readily ascertained when the sequence of the full cDNA insert is determined in accordance with Example 13.

Example 4

EST Sequences Characterization: Second Set

The ESTs of Example 2, including SEQ ID NOs 316-2407, were screened against known sequences listed in GenBank and other databases, as in Example 3. The results are reported in Table 2. The quality of the match is given as percent identity and length in base pairs for nucleotide matches and amino acid residues for peptide matches. In many cases ESTs match multiple domains on several related proteins; for example, EST00825 matches two transmembrane domains on both GABA and Norepinephrine transporters. Nucleotide databases are: GenBank (GB), and EMBL (E); peptide databases are: GenPept (GPU), Swiss-Prot (SP), and PIR.

The great majority (83%) of the partial cDNA sequences reported in Example 2 are unrelated to any sequences previously described in the literature. Based on database matches to known genes from humans as well as from such evolutionarily distant organisms as E. coli, yeast, elegans, Drosophila, barley, Arabidopsis, rice, and green algae, we have preliminarily identified the functional type of a number of the ESTs (Table 2). These include a novel gene similar to Notch/Tan-1 (Adams et al., supra), a new neurotransmitter transporter gene, and a new member of the multi-drug resistance gene family. Several genes involved in development or cell differentiation in Drosophila are represented by cimilar human ESTs, including seven in absentia (Carthew, R. & Rubin, G. Cell 63: 561-577-(1990)),

big-brain (bib) (Rao, Y., Jan, L., & Jan, Y. Nature 345: 163-167 (1990)), the discs tumor suppressor (Woods, D. & Bryant, P. Cell 66: 1-20 (1991)), and the homeotic gene orthodenticle (Finkelstein, R., Smouse, D. Capaci, T., Spradling, A. & Perrimon, N. Genes. Dev. 4: 1516-1527 (1990)). New members of gene families previously known in humans include a Ca⁺²-transporting ATPase, an ADP ribosylation factor, and a new neural-cell adhesion molecule gene.

The 1971 ESTs without a putative identification were analyzed using the coding-region prediction program CRM via the GRAIL server (Uberbacher, E. & Mural, R. Proc. Natl. Acad. Sci. USA 88: 11261-5 (1991)). Fifteen percent of the unknown ESTs scored an excellent probability of containing protein-coding sequence. Fifty percent of the ESTs to known human genes contain protein-coding sequences, therefore, at most half of the unknown ESTs are likely to contain coding sequences. We have found no evidence that genomic DNA or cDNA to unspliced precursor RNA is a major contaminant of either the hippocampus or fetal brain library.

Table 2: ESTs Identified by Database Matches

SEQ ID EST# Putative Identification	Accession DB Len %ID
208 EST00250 60K filarial antigen	A28209 PIR 108 56.9
2320 EST01784 60K filarial antigen	A28209 PIR 88 50.6
969 EST01982 ADP-ribosylation factor 1	B33283 PIR 84 41,2
1834 ESTO1620 AMP deaminase, brain	A37056 PIR 57 100.0
97 EST00289 Aconitase	A35544 PIR 105 90.6
251 EST00370 Actin, other	S10021 PIR 44 51.1
248 EST00271 Actinin, alpha	HUMACTAR GB 271 85.3
891 ESTO1891 Actinin, alpha	HUMACTAR GB 315 81.6
1500 EST02538 Actinin, alpha	HUMACTAR GB 271 75.0
132 EST00110 Agrin	RATAGR GB 269 82.2
1852 EST01625 Agrin	RATAGR GB 103 84.6
. •	HUMALA GB 92 82.8
691 EST00675 Alcohol dehydrogenase	RICGOS2G_1 GPU 38 59.0
2408 EST00244 Amyloid A4	HUMAFPA4 GB 135 91.9
1965 EST01664 Amyloid A4	A29030 PIR 52 54.7
2068 EST01694 Amyloid A4	QRHUA4 PIR 83 69.0
2092 EST01700 Anion exchanger homolog AE3	A33638 PIR 95 97.9
2092 EST01700 Anion exchanger homolog AE3 1880 EST01634 Axonal glycoprotein TAG-1	A34695 PIR 69 87.1
1492 EST02530 B cell-specific Mo-MLV integration site	1 (bmi-1) MUSBMI1A GB 111 87.5
1277 ESTO2306 Bib protein	S09699 PIR 57 53.4
13 EST00255 Cadherins	CADN\$HUMAN SP 41 45.2
1348 EST02378 cAMP-dependent protein kinase inhibito	
1931 ESTO1041 cAMP-regulated phosphoprotein	B35308 PIR 21 86.4
1413 EST02447 cAMP-specific phosphodiesterase	HUMPDEAA GB 363 69.0
396 EST01443 CDPdiacylglycerol-serine O-phosphatidyl	transferase JH0368 PIR 33 41.2
396 EST01443 CDPdiacylglycerol-serine O-phosphatidyl 1956 EST01663 Ca2+-transporting ATPase 2 1126 EST02146 Calbindin D28	B28065 PIR 125 88.9
1126 EST02146 Calbindin D28	RATCALBD28 GB 81 87.8
1039 EST02055 Calcium channel	S05054 PIR 33 67.6
1910 EST01645 Calmodulin	RATRCM1 GB 120 90.1
485 EST01466 Calmodulin-dependent protein kinase, ty	rpe II, beta A26464 PIR 93 98.9
913 ESTO1913 Clathrin coat assembly protein AP50 ho	molog YSCYAP54_1 GPU 62 63.5
	PIGCOFIL GB 132 89.5
2400 ESTO1824 Cysteine-rich intestinal protein	GYRTI PIR 56 66.7
1588 EST02633 D22Z3 repetitive DNA	HUMREP GB 160 76.4
2192 ESTO1257 Diacylglycerol kinase, lymphyocyte	S09156 PIR 44 42.2
1441 EST02477 Diamine acetyltransferase	ATDASHUMAN SP 74 45.3
650 EST00642 Dilute (myosin heavy chain)	MUSDILUTE_1 GPU 27 100.0
2302 EST01779 Discs-large tumor suppressor	DRODLGA_1 GPU 53 63.0
188 EST00256 Enhancer of split	A30047 PIR 86 58.6
2289 ESTO1325 Fatty acid synthase	RATFAS GB 98 79.8
310 EST00377 Fo ATPase beta subunit, mitochondrial	BOVMTASB GB 293 85.4
1332 EST02362 GA binding protein, beta subunit	MUSGAC_1 GPU 86 90.8
1667 EST00825 Gamma-aminobutyric acid transporter	A35918 PIR 26 59.3
2217 EST01738 Gelation factor ABP-280	A37098 PIR 74 80.0
1412 EST02446 Glutamate-aspartate carrier protein	JV0092 PIR 57 37.9
1020 EST02034 Glutaminase	GLS\$RAT SP 34 74.3
1885 EST01639 Histocompatibility antigen modifier 1	A37779 PIR 63 75.0
1495 EST02533 Hypothetical 43.5K protein	JU0319 PIR 43 52.3
2326 EST01791 Inositol-1,4,5-trisphosphate 3-kinase	JN0129 PIR 65 68.2
SEQ ID EST# Putative Identification	

724 EST01529 Interferon-induced 54K protein	INI4\$HUMAN SP 76 70.1
1035 EST02051 J1 protein	MUSJ1PRO GB 362 85.7
1229 EST02258 KUP protein	HUMKUPMR 1 GPU 54 36.4
993 EST02007 Kinase 5 protein	CHKCEK5_1 GPU 68 94.2
77 EST00257 Kinesin	A35075 PIR 57 86.2
78 EST00258 Kinesin	A35075 PIR 62 47.6
2245 EST01748 Kinesin	A35075 PIR 98 52.5
2282 EST01764 Lamin B receptor	A36427 PIR 76 71.4
2173 EST01724 Lon protesse	JQ0901 PIR 103 41.3
1427 EST02463 Long-chain-fatty-acid-CoA ligase	A36275 PIR 36 62.2
313 EST00276 Lysosomal membrane glycoprotein 1 (L	
161 EST00247 MARCKS (myristoylated alanine-rich pro	
1386 EST02418 MARCKS homolog	MMF52 EU 237 92.4
769 EST00734 MARCKS homolog	S08341 PIR 61 40.3
43 EST00371 Maternal G10 protein	S05955 PIR 38 92.3
1468 EST02505 Matrin 3	RATMATRIN3 GB 137 93.5
639 EST00632 Membrane transport superfamily (GTP-d	
1894 EST01643 Membrane transport superfamily (GTP-	•
824 EST01865 Microtubule-associated protein 1B	RATNEU GB 293 86.4
223 EST00368 Microtubule-associated protein 1B	A33645 PIR 30 54.8
2032 EST01683 Microtubule-associated protein 18	A33645 PIR 49 62.0
2017 EST01678 Milk fat globule membrane protein	A36479 PIR 48 61.2
1704 EST01580 Myeloid differentiation primary response	
2226 EST01744 NAD(P) + transhydrogenase (B-specific)	
1567 EST02610 Neural cell adhesion molecule L1	S05479 PIR 82 43.4
506 EST01471 Neuraxin	S06017 PIR 120 84.3
1566 EST02609 Neutrophil oxidase factor	A34855 PIR 43 47.7
952 EST01961 Notch/Xatch	HUMTAN1 1 GPU 85 57.0
227 EST00259 Notch/Xotch	A35844 PIR 74 85.3
1395 EST02429 Nuclear factor 1-like protein (NF1)	HAMNF1A GB 111 92.0
1881 EST01573 Nucleoside diphosphate kinase	A33386 PIR 71 52.8
346 EST01828 Otd homeotic protein	A35912 PIR 35 52.8
. 2254 EST01751 Phosphatidylinositol-4,5-bisphosphate p	phosphodiest A28807 PIR 40 90.2
1869 EST00992 Polymyxin B resistance	A32714 PIR 20 76.2
93 EST00287 Processing enhancing protein	S03968 PIR 96 58.8
2353 EST01806 Prohibitin	RATPROHIB_1 GPU 120 97.5
2297 EST01775 Prohormone cleavage enzyme	MUSMPC1A_1 GPU 91 93.5
9 EST00376 Prolyl endopeptidase	PIGPREP GB 223 83.9
1089 EST02087 Protein kinase C, zeta	HUMPKCL GB 382 58.7
1933 EST01650 Protein phosphatase 2A beta subunit	HUMPROP2AB GB 288 76.8
202 EST00298 Protein-tyrosine phosphatase LRP	LRP\$MOUSE SP 62 44.4
1654 ESTO1572 Protochlorophyllide reductase	S04783 PIR 34 57.1
38 EST00374 RNA polymerase II 6th subunit (RPO26)	
1478 EST02515 Rab5	F34323 PIR 91 82.6
2368 ESTO1389 Radial spoke protein 3	S05962 PIR 58 52.5
37 EST00038 ras p21-like small GTP-binding protein (s	
180 EST00299 ras-related proteins	S10493 PIR 51 46.1
1700 EST01579 Retrovirus-related gag polyprotein	FOHUE2 PIR 95 77.1
1511 ESTO2550 Retrovirus-related pol polyprotein	GNLJGL PIR 50 54.9
102 EST00248 rho H12/ ARH12	BOVBGBRH GB 195 79.6
1715 EST01583 Ribosomal protein L18a	R5RT18 PIR 68 95.7
SEQ ID EST# Putative Identification	Accession DB Len %ID

Park to make a processing the second

```
1856 ESTO1627 Ribosomal protein L1a
1974 EST01667 Ribosomal protein L3
                                                   JQ0771
                                                               PIR 74 80.0
 301 EST00300 Ribosomal protein L30
                                                   R6RT30
                                                               PIR 57 96.5
 22 EST00301 Ribosomal protein S10
                                                   R3RT10
                                                              PIR 66 97.0
2402 EST01826 Ribosomal protein S10
                                                    R3YM10
                                                               PIR 36 51.4
 463 EST01459 Ribosomal protein YL10
                                                    S11581
                                                               PIR 40 68.3
                                                              PIR 46 80.8
PIR 97 77.5
1408 EST02442 Seven in absentia
                                                  A36195
 299 EST00249 smg p25A GDP dissociation inhibitor
                                                      A35652
 951 ESTO1960 Spectrin, beta
                                                 HUMSPTB GB 268 67.7
2089 EST01699 Sperm membrane protein
                                                               PIR 52 58.5
                                                     A35981
2073 EST01697 Succinate dehydrogenase flavoprotein
                                                      BOVSDHFP1_1 GPU 44 100.0
2138 EST01715 Succinate dehydrogenase flavoprotein
                                                       BOVSDHFP1_1 GPU 49 92.0
430 EST00472 Synaptotagmin (p65)
                                                   SY65$HUMAN SP 27 53.6
1371 EST02402 Talin
                                               MUSTALINR_1 GPU 79 81.2
1771 EST01601 Thiosulfate sulfurtransferase (rhodanese)
                                                     ROBO
                                                                PIR 65 81.8
 300 EST00232 Transforming protein (dbl)
                                                   TVHUDB
                                                               PIR 25 65.4
                                                         PIR 33 67.6
 189 EST00282 trkB
                                               A35104
653 ESTO1512 Tubulin, alpha
                                                HUMTUBAG
                                                             GB 223 75.0
 594 ESTO1490 Tubulin, beta
                                                 HUMTBB5
                                                             GB 298 93.6
757 EST01542 Tubulin, beta
                                                 HUMTUBBM
                                                             GB 217 90.4
1245 ESTO2274 Tubulin, beta
                                                 A26561
                                                            PIR 105 88.7
1147 EST02169 Tyrosine kinase
                                                 HUMECK
                                                             GB 384 74.3
1701 EST00853 Unc:104
                                                            NR 36 45.0
2121 EST01711 Valine-tRNA ligase
                                                  A29871
                                                             PIR 56 57.9
187 EST00152 Wilm's tumor-related protein
                                                  HUMQM
                                                                GB 228 99.6
1726 EST01588 XPR2 alkaline extracellular protease
                                                    826955
                                                                PIR 88, 46,1
249 EST00275 Zinc Finger Proteins
                                                             PIR 25 57.7
                                                 S06551
413 EST01446 Zinc Finger Proteins
                                                 S00754
                                                             PIR
                                                                  45 60.9
469 ESTO1460 Zinc Finger Proteins
                                                 C32891
                                                             PIR
                                                                 34 54.3
833 EST01560 Zinc Finger Proteins
                                                  S00754
                                                             PIR 105 67.0
1230 EST02259 Zinc finger proteins
                                                  S00754
                                                             PIR
                                                                 71 62.5
1496 EST02534 Zinc finger proteins
                                                  A34612
                                                                 50 . 45.1
2324 EST01352 Zinc Finger Proteins
                                                  S10397
                                                             PIR'
                                                                 29 56.7
```

10

15

20

25

There is little redundancy in EST sequencing according to the present invention. Of the nuclear-encoded messenger RNAs, the most common ESTs were to the β -actin (0.6% of the EST clones) and myelin basic protein genes (MBP, 0.5% of the clones). MBP, a highly expressed structural component of nerve tissue (Kamholtz, J., de Ferra, F., Puckett, C., & Lazzarini, R. **Proc. Natl. Acad. Sci., USA 83**: 4962-4966 (1986)), displays four alternate splicing forms, of which at least two are present among the ESTs reported here. Other common ESTs were Gs-alpha gamma-actin and both a- and alpha-tubulin.

By matching ESTs to known database sequences, a phenotypic characterization of the tissue begins to emerge. Protein superfamilies matched by ESTs were grouped into three broad functional categories to assess the biological spectrum represented by these randomly selected cDNA clones. Structural and metabolic classes comprised about 30% of the ESTs with database matches. Twenty-five percent were involved in regulatory pathways and the remainder were not classifiable. Eleven of the eighteen enzymes of glycolysis and the citric acid cycle are represented by at least one subunit or isozyme. In addition, several genes not previously known to be expressed in the brain were matched, including spermine/spermidine acetyltransferase (Casero, R., Celano, P, Ervin, S., Applegren, N., Wiest, L. & Pegg, A. J. Biol. Chem. 266: 810-814 (1991)) and osteopontin (Young, M., Kerr, J., Termine, J., Wewer, U., Wang, M., McBride, W. & Fisher, L. Genomics 7:491-502 (1990)).

30

EXAMPLE 5

Mapping of ESTs to Human Chromosomes

Randomly selected ESTs corresponding to SEQ ID NOs. were assigned to chromosomes via PCR (see Table 3).

Oligonucleotide primer pairs were designed from EST

10

20

25

30

5 د

sequences to minimize the chance of amplifying through an intron. The oligonucleotides were 18-23 bp in length and designed for PCR amplification using the computer program INTRON (National Institutes of Mental Health, Bethesda, MD). The program is based on the assumptions that: 1) introns are genomic sequences that interrupt the coding and noncoding sequences of genes (Smith, J. Mol. Evol. 27:45-55 (1988)); 2) there are consensus sequences for splice junctions (Shapiro, et al., Nucl. Acids Res. 15:7155-7174 (1987)); and 3) that 90% of the human genes studied have 3' untranslated regions of mRNA not interrupted by introns in the genomic DNA (Hawkins, Nucl. Acids Res. 16:9893-9908 (1988)).

The program evaluates the likelihood that a given GG or CC dinucleotide represents a former exon-intron boundary. Specifically, every input strand is processed by the INTRON program twice, first evaluating the sense mRNA strand, and then processing the complementary or anti-sense strand. The program evaluates each sequence by finding all GG or CC pairs (possible former splice sites), searching for STOP codons in all three reading frames, and analyzing the GG or CC pairs surrounded by stop codons. All regions of the EST that are unlikely to contain splice junctions based on CC content, GG content, and stop codon frequency are then marked by the program in uppercase.

The creation of PCR primers from known sequences is well known to those with skill in the art. For a review of PCR technology see Erlich, H.A., PCR Technology; Principles and Applications for DNA Amplification. 1992. W.H. Freeman and Co., New York. ESTs were examined for the presence of stop codons in each reading frame and for consensus splice junctions. The presence of stop codons and absence of splice junction sequences are more characteristic of 3' untranslated sequences than of introns. The untranslated sequences are unique to a given gene; thus, primers from

10

15

20

25

30

35

these regions are less likely to prime other members of a gene family or pseudogenes.

The primers were used in polymerase chain reactions (PCR) to amplify templates from total human genomic DNA. PCR conditions were as follows: 60 ng of genomic DNA was used as a template for PCR with 80 ng of each oligonucleotide primer, 0.6 unit of Tag polymerase, and 1 uCu of a ³²P-labeled deoxycytidine triphosphate. was performed in a microplate thermocycler (Techne) under the following conditions: 30 cycles of 94°C, 1.4 min; 55°C, 2 min; and 72°C, 2 min; with a final extension at 72°C for The amplified products were analyzed on a 6% polyacrylamide sequencing gel and visualized by autoradiography. If the size of the resulting product was . equivalent to the EST from which the primers are derived, then the PCR reaction was repeated with DNA templates from two panels of human-rodent somatic cell hybrids; BIOS PCRable DNA (BIOS Corporation) and NIGMS Human-Rodent Somatic Cell Hybrid Mapping Panel Number 1 (NIGMS, Camden, NJ).

PCR was used to screen a series of somatic cell hybrid cell lines containing defined sets of human chromosomes for the presence of a given EST. DNA was isolated from the somatic hybrids and used as starting templates for PCR reactions using the primer pairs from EST sequences selected above. Only those somatic cell hybrids with chromosomes containing the human gene corresponding to the EST will yield an amplified fragment. ESTs were assigned to a chromosome by analysis of the segregation pattern of PCR products from hybrid DNA templates. For a review of techniques and analysis of results from somatic cell gene mapping experiments. (See Ledbetter et al., Genomics 6:475-481 (1990).) The single human chromosome present in all cell hybrids that give rise to an amplified fragment represents the chromosome containing that EST.

WO 93/16178 PCT/US93/01294

-48

The assignment of 100 ESTs and corresponding genes to chromosomes by PCR is shown in Table 3.

24.77.

Table 3: Assignment of ESTs to Chromosomes by PCR

			-	of con
SEO ID	EST#	Chi	PRIMER #1	PRIMER #2
5	EST00012	1	TCCAGGCAATCCCAGAATAG	CTAATTGAGCTCACTGGCCC
57	EST00058	1	CTGTTTGCAAGTTTCAAAGC	GCCATTTCTAACAACCAGAG
. 64	EST00066	1	GCCATTGTGCTGAATAGAGT	GTTAGTGTTTCCTTAGCAAG
83	EST00079	1	CAGCTAATTGACCTGGGCTA	CAACATGCTCTGAGCTTTAG
83	EST00079	1	GGCAGAGCATAATGAGTATA	CATATGCATATGGTCCCTAT
91	EST00086	1	AGTTTAGATGGAGGGCTGTC	TCTGCCCTAATGCGCAGGCT
105	EST00365	ī	CTTAATCACCTCCCTTTTGT	
109	EST00095	1	AGTCTAATCCTGTACACTTG	CCCCCTTTGTGTGTAAGGTC
116	EST00100	ī	TTAGAAGTGCCCATGGGAGG	CGGGCTTTCTCTGAATTGGT
141	EST00118	ī	CTCAGAGAAACTTAGGTGAA	TTTTAAGGCTCTGGAGTGTT
220	EST00372	ī	AAGTTGCACATTGCCCAAGG	CTACAGAATCATTTCACCAG
237	EST00187	ī	TTACAAATTTCTCTTGACGC	ATAGTACTGCAAGGTTATTC
242	EST00192	ī	GGATCAGATAATCAAACAGG	CTGAAGGAGCACAGTTTCTC
259	EST00202	î	GCATCACAGTTTAACTGAGG	GCTTAGGATATGAATGCATA
269	EST00293	ī	CTGTTGCTGTGCAGTAGCTT	CTACATATTTGTGCCTCCTT
299	EST00249	ī	GATCATGCAGACGTAGATAT	CTTTTGACCCAGTGAAACTT
1651	EST00249	ī	TAGTCGCTGTAAGTTGATTC	CCAACTCCTGCCAGATCATT
16	EST00010	2	CAGGCAAGTTTCTTCCAGGA	GCTTTGCTGGATGCTTCATT
1898	EST01013	2.	GGCTGAGAACGGTTAGCATA	TCAGACCCATGGTCAGCTT
8	EST00234	2	TAGAAGGCAAACTATGTCCC	CCCTCAGCTTAGGGGAATG
36	EST00234	2	AGCCAGAAGGCTGCTTAAAG	GGTTGAGGATTGGCTTTTAC
123	EST00037	2	GTCTAATTTGTAACCTTCAG	GCAGTGAACCAGTACTCCTA
192	EST00100	2	GATTTATGTCTGGGAACTAA	GATAGATTGTATAAGAAGCC
200	EST00162	2	TTTAATGGGTGGTGGGAGCT	GCAGCATGTGAAAGAATGAT
284	EST00162 EST00216	2	CCTAAGAATTCGTTTGGCTC	CGATGCACATCCTTCTCCAT
102	EST00218	3	ATACTACATCTAGTCTGG	GTCTGGCACATAATAGATTTG
167	EST00248	3	AAACAGCTGCGGAGTACA	TTACAGTTCTGTGGTTTC
12	EST00136	3	CCTAGCAAACTCATACACAC	AAAGGATCCTCCACTCCAGA
60	EST00062	3	ACACATTAACGGTGCTGCAG	CATAAGTGAATGGACACAGG
77	EST00052	3	AAGCTCACAACGCAGATCTG	GGAATCAGCCCTTGAGGACT
107	EST00257	3	ATTGAACTCTGTCAACAGTG	CTGGAACAGCTTACAAAGGT
108	EST00094	3	AL2-GCAGGATGTCAGTCTTTTGAG	TGTAAAACAAAGGCCAAACT
1706	EST00857	3	AL2-GCAGGATGTCAGTCTTTTGAG	AGCACACATTATCTACCACGGC
37	EST00037	3 4	AACTTCGCAGTCATGAGAAC	CCAGCACACATTATCTACCACG
6	EST00038	4	CACATGTTCTCCCTCTTTCA	TGTATCGGGCAGTTCTCAG
37	EST00013	4	AL2-GGAAGTACAGGATTTGGC	GCATTTTGGAGCTCTTCCGT
31	EST00038	5	TGGGTACCCTAAGGTGTTTG	TTAGAGATGGGATGATGCCG
28	EST00033	5		GACTAATCTAAGGTCTAGG
59	EST00030		AGATAAGTTAGGAAGCTGGT	ACTCACTGCTAGTATCATCC
74	EST00061	5 5	AAAGTTTCTTAGCACCCCCC	CAGACTTTGACAAAAGAATC
121	EST00073 EST00104		ATCAGACACGTGGCAGGGTT	AAGTCCCTGAGGGTGCAGAA
149	EST00104 EST00123	5	TGAAGGCAGCTGCTAAATCT	GGATGTATTGATCTGACTCA
235	EST00123 EST00185	5	ATACTGTCAACGGAGGGTGA	GTCTGCAGGTTTCTCCTTGA
1643	EST00185	5 5	TTACTGTCCCATCAGATATC	TACACTCTTAAGAAGGTATG
1677			GAGCGTTTAAAAGAGATTCT	TACAGACAGCCATGTTCCAA
23	ESTO0835	5 5	AL2-TCTCCAACACAGTCATGC	CGGATGCCATCATATACC
121	EST00026 EST00104	5	CCTGCAGTGACACTTAACAT	CTGCTCACCTGAAATTGATAC
461	23100104	5	AL2-CAGATCAATACATCCTCTGGG	CTGTGCAGTGGTGAGTAAAAGG

			•	• .
SEO ID	EST#	Chr	PRIMER #1	PRIMER #2
1	EST00007	6	TAGTTGATGGTCTGGGTTAT	GAAATCCCAGGGAGACAATG
19	EST00023	6	CAACTTACATTAGGGGTTTG	GACCTCATTAGAAGAGCCCA
155	EST00129	6	GGAAGCTGCCATATAAGCTC	TCAGTGTCGTACAATCTACC
224	EST00356	6	GCTGTATGTTAACCCTTTGT	TGGAACCCTCAAACACTGCT
288	EST00219	6	ACTTTCATGTTGAGAAGTAT	ATCTAGCTGAAACATTGCTG
1638	EST00798	6	CTTCATCTGTTAACTGTTGA	TGAAAATGAGTCACAGGCAG
1675	EST00833	6	AL2-ACCCAGTTCTCAAAGACC	GGTTTACCATTCAGAGGC
22	EST00301	6	CTCCGTGATTACCTTCATCT	TTGTAGGTATCTCTGTCAGCT
207	EST00167	7	GGTGCTACTTTGTGAATGCT	AGCAATGTGATTTTGTAGG
137	EST00272	. 7	AGTGGTCACTATCTACATGG	GATTCAGAATTACTAAGCCG
1659	EST00817	7	TGTATAGGCTCTACATAAAG	CTTAATCATGGATTCTTCGT
1680	EST00838	7	AL2-GTTCTTTCCCAGGTATGC	TTGTTGGTACTGAGGAAGTGCG
292	EST00223	8	TGCAGCAGTGACCATGAGAA	ATCATCTTTCCACGCGCCTT
134	EST00225	9	TCTGGGCTTCTGTGGTTCAA	CTGGCTGCTCAGCAACTCAT
1906	EST01021	9	GGATGTTTTCTATGTGACGA	TTCCAGTGCCCCTTTTGTCC
1645	EST00804	10	CTCCTTTGGGACAAACAACT	CCAACCCAAACATATTCTA
20	EST00024	10	AGCTGTTCCTGAGAGATGCA	CCTTGTGAAGAAGACTTTC
157	EST00024	10	TCAGCAACAGGTCACTTTGG	
172	EST00131	10	TACTAGCATTTCTTACTCTC	CTAAGCATCTGCATGTCCAG TATGCTGATTGTTTGCACTC
250	EST00142 EST00197	10	GGTGATTAGAGAGTCTGTTG	
133	EST00197	11	GGAAATTAGGCTTAGCTCAC	GAACTCTGTAGTGTTCTAAA
178	EST00111 EST00294			GTGCAGAATACTTAGAGTCC
10		11 11	GTTTGAAGGAAGTGATTTCC	TAGGGCCACCTCCAGTTCAT
	EST00016		GTCTTTGGATTCTACGTAGA	CGATAATGACATTTCTTCTGG
126 7	EST00109 EST00014	11 12	AL2-CTAACCACACCCACACATTG AACTTGCAACATAAATACTAG	CCTCAGCACAAGAGAAGAATGG
254				GAGCAATGATTTCTAACAGT
2409	EST00200 EST00273	13	TTGTGTACTGTCTGATAGAC	TAAGCCATGGGCATCTATAA
170	EST00273	13 14	GCAAGATGATGGAACATCCC GGTGCTTAAGGCCACTTTTG	TTCCTTCTGGAGGCTCTACA
255	EST00295 EST00201	14	,	CTTAGAGGATCATAGGTCTG
290	EST00201		CCAGGAGAGTAAGAAGATCA GTGCCAAGATGGCTCATGTA	GCAGAGTTGAATATGAACCT
293		14		GTATAGCTTTAAGCCAGTTC
	EST00224	14	AATGCATTATGCCTGGTCTT	GGAAAAGTCTAGAACTTAGT
1664	EST00822	14	GGGTCAGAATTAAGAGGTCT	GTTCATCTCTAACTCCTTTC
:315 1689	EST00008	14	AAGCTGGCTGGGAAATGTTC	GTCATGCTAGTAAACTTACAC
	EST00845	14	AL2-AGGAGGAAGCTGAAATCC	GGAAGTCCATAAGAGACTCACC
95	EST00088	15	GTGACAGACCATGTCTATTG	AAGTGAGCGATTGCACCTTC
205	EST00165	15	AGGATGACCTGAGTGAGCTG	CCATGGCAGCAAGGAACTCT
33 247	EST00034	16	TGTGTGAAAGGGAGTCTTGT	CCATTTTGACTGTTCCATAG
	EST00279	16	TGGCTAGGGCAGGCCTTAAA	GAGAAGAATATCAAATGGGG
18	EST00373	16	CCATCTGTGTCCCAATTAAGC	AGGGAAGAAGTCTAGAGCGA
68	EST00068	17	CAAAGACGGGAGACGAATGA	AGTGGAACGCGTGGCCTATG
1652	EST00811	17	GAGCTGCATGTTGATAAGTA	TTGACTTAAGCTGACCTTAA
1702	EST00854	17	AL2-TTGCTGTGGAATCCATGAGAG	GGCAAGTGATCTGTTCTTGG
84	EST00080	19	AGAGATGTCAGTCCATTATC	CTATTCCACCTTACTCAAGG
223	EST00368	19	CATCATGTCGGAGACGCATT	TGGATGACCTGAGTCTGCAG
21	EST00025	20	AGTTCTGGAGGCTAGGAGTT	ATGTAAGGACCCCTAGATGG
210	EST00168	20	TGTCAACTTCCCTTTGGCCT	GAAGCTTGCTCATTCAGGAA
136	EST00113	20	AL2-TCGGAGAAGTTGCAGTTTCTG	GTTAAAAGCTGTTAGACGGGGC
120	EST00103	22	CACTGACTGACTCCTCTTTA	GGAACCGTAACTCTCCATAG
313	EST00276	Х	ATTGACCTTCAATGTAATAA	TTGGATTGGGCAAAATAG

-51-

SEO ID	EST#	<u>Chr</u>	PRIMER #1	PRIMER #2
162 1669 1917 1708	EST00133 EST00827 EST01029 EST00858	X X X	ATGTGAGCATCTATACCTGC CGGACAACTAGGATAAATGC GAATAGCATTATTAGCCAGT AL2-AAGGCGAGGATTATGTGC	AATGAAGGCATGAGAATAGG TACGCGTTTGAATGGCTTGA GGACCTATTGGAGATCTACT TTCTACTGGGTACACTTCGACC

Abbreviation: AL2: Amino-Link-2 Fluorescent Tag, Chr.: Chromosome.

The foregoing techniques have been used to further localize 9 ESTs and their associated genes to precise locations onto chromosome 6 or chromosome X, as reflected in Table 4A (in Example 7 below), using sublocalization techniques that employ somatic cell hybrids. used as hybridization probes and mapped to other chromosomes using techniques disclosed in Example 7. Somatic cell hybrids were prepared that contained defined subsets of chromosomes 6 and X. Methods for preparing and selecting somatic cell hybrids are known in the art. review of an exemplary procedure to generate somatic cell hybrids containing the short arm of human chromosome 6, see Zoghbi, et al., Genomics 9(4):713-720 (1991). general review of somatic cell hybridization see Ledbetter et al.(supra). The hybrids were processed to obtain DNA and analyzed by PCR and by fluorescence in situ hybridization. SEQ ID NOs 19, 22, 1, 224, 288 mapped to chromosome 6, while SEQ ID NOs 162, 1917, 1699 and 1899 mapped to chromosome X using somatic cell hybrids.

20

10

15

EXAMPLE 6

Mapping of All ESTs to Human Chromosomes

The procedure of Example 5 is repeated for all of the ESTs from Examples 1 and 2 not previously mapped to human chromosomes. Data are generated corresponding to the data in Table 3 for all of the unmapped ESTs. As previously mentioned, virtually all of the ESTs will map to a unique chromosomal location. The inability of any ESTs to localize to a unique location will be readily ascertainable during the mapping process.

30

25

Physical mapping of the type reported in Table 4 on all the EST clones reported here would provide human chromosome markers spaced on average every 1.2 megabases and would roughly double the number of expressed sequences that have been localized to chromosomes (McKusick, V. FASEB

WO 93/16178 PCT/US93/01294

-53-

J. 5: 12-20 (1991)). Mapped ESTs are also a new resource to identify candidates for the estimated 5000 single-locus disease-associated genes (Id.).

EXAMPLE 7

Alternative Technique for Mapping to Chromosomes Mapping of ESTs to chromosomes using fluorescence in situ hybridization

5

10

15

20

25

30

This technique was used to map an EST to a particular location on a given chromosome. Cell cultures, tissue, or whole blood were used to obtain chromosomes.

0.5 ml. of whole blood was added to RPMI 1640 and incubated 96 hours in a 5%CO₂/37°C incubator. 0.05 ug/ml colcemide was added to the culture one hour before harvest. Cells were collected and washed in PBS. The suspension was incubated with a hypotonic solution of KCl added dropwise to reach a final volume of 5 ml. The cells were spun down and fixed by resuspending the cells in methanol and glacial acetic acid (3:1). The cell suspension was dropped onto glass slides and dried.

The slides were treated with RNase A and washed then dehydrated in a series of increasing concentrations of ethanol.

The EST to be localized was nick-translated using fluorescently labeled nucleotide (Korenberg, Jr., et al., Cell 53(3):391-400 (1988)). Following nick translation, unincorporated label was removed by spin dialysis through Sepharose. The probe was further extracted with phenolchloroform to remove additional protein. The chromosomes were denatured in formamide using techniques known in the art and the denatured probe was added to the slides. Following hybridization, the cells were washed. The slides were studied under a fluorescent microscope. In addition, the chromosomes can be stained for G-banding or Q-banding using techniques known in the art. 24. TT

30

35

The resulting metaphase chromosomes had fluorescent tags localized to those regions of the chromosome that were homologous to the EST. Thus, a particular EST was localized to a particular region on a given chromosome. In this manner, SEQ ID NOS 396, 485, 506, 1880 and 1894 were mapped using fluorescent in situ hybridization to locations on chromosomes 17, 7, 10 and 1 respectively (See Table 4B below). For a review of the technique see Verma et al., Human Chromosomes: A Manual of Basic Techniques. Pergamon Press, NY (1988), which is hereby incorporated by reference.

Table 4: Precise Chromosomal Localization of ESTs

•		SEQ ID	EST#	Map Location
	A.	19	EST00023	6p
15	•	22	EST00301	6p
	n	1894	EST01643	6p21
		1	EST00007	Ga Car
		224	EST00356	6q
		288	EST00219	6q
20		162	EST00133	Xp11.21 - Xp21.2
		1917	EST01029	Xp11.21 - Xp21.2
		1669	EST00827	Xq26 - Xq27.1
,		1899	EST01014	Xq28
	В.	1880	EST01634	1q32
25		485	EST01466	7p13
		506	EST01471	10q11.2
		.396	EST01443	17q25

EXAMPLE 8

Automated DNA Sequencing Accuracy

ESTs that match human sequences in GenBank are excellent tools for the analysis of the accuracy of double-strand automated DNA sequencing. Ninety EST/GenBank matches were examined for the number of nucleotide mismatches and gaps required to achieve optimal alignment by the Genetics Computer Group (GCG) program RESTFIT (Devereux et al, Nucleic Acids Research 12: 387 (1984)).

The number of mismatches, insertions and deletions was counted for each hundred bases of the sequence (Table 5). As expected, the sequence quality was best closest to the primer and decreased rapidly after about 400 bases. The number of deletions and insertions relative to the GenBank reference sequence increased five- to ten-fold beyond 400 bases, while the number of mismatches doubled. The average accuracy rate for individual double-stranded sequencing runs was 97.7% to 400 bases.

्रः । इक्रायम्

TABLE 5. Accuracy Of Single-Run Double-Stranded Automated Sequencing

Bases from Primer	Mismatches/	Gaps	Percent	Aligne	d
	Ambiquities ⁺	<u>Insertions</u> [†]	Deletions	Accurate	Bases
101 - 200	1.45	0.18	0.19	98.2	8,800
201 - 300	1.72	0.25	0.11	97.9	8,130
301 - 400	2.07	0.98	0.37	96.6	5,404
>400	3.53	2.63	1.06	92.8	3,197

ESTs statistically identical to known human sequences and those matching mitochondrial and ribosomal genes were aligned with sequenced from GenBank using the GCG program BESTFIT. The first 85 nucleotides was polylinker sequence which was not aligned with the pBluescript SK reference sequence. Tabulation of errors began 15 bases into the BESTFIT alignment and thus is reported beginning with bases 101-200. *Error rates are reported as number of mismatches, insertions, or deletions per hundred aligned bases. "Mismatches" includes ambiguous base calls.

10

15

44.44

EXAMPLE 9

Probability of ESTs Containing Coding Sequences

The ESTs of the present invention were statistically evaluated using the coding-region prediction program CRM via the GRAIL server (Uberbacher, E. & Mural, R. Proc. Natl. Acad. Sci. USA, 88: 11261-5 (1991)). The CRM program uses a neural network to combine results from several different coding regions by looking at different 6 bp sequences found in coding exons and in introns. program additionally conducts reading frame searches and assesses randomness at the third position of codons. protocol categorizes sequences as having an excellent, good, marginal, or poor probability of containing coding The results are reported in Tables 6-9. There were 219 ESTs categorized as "excellent" (Table 6); 120 categorized as "good" (Table 7); 113 categorized as "marginal" (Table 8); and 1743 categorized as "poor" (Table These results indicate that most ESTs of the present invention comprise noncoding regions.

Table 6: ESTs with Excellent Probability of Containing Coding Sequence

•							
SEQ ID#	EST#	973	EST01987	4007			·
<u> </u>	2017		EST01993	1807			2373 EST01393
. 7	EST00014	ÓRÓ	EST01994	1809			2374 EST01394
15	EST00020	984	EST02000	1820	EST00951		2393 EST01417
48		1000	EST02014	1829			2394 EST01418
				1849			2396 EST01420
62		1004		1860			
66		1007		1866	EST00989	•	
75	EST00074	1018	EST02032	1871	EST00994	•	
· 98	EST00260	1021	EST02035	1888			
106	EST00092	1034	EST02050	1890	EST01007		
108	EST00094	1047	EST02063	1892			
114	EST00098	1000	EST02109	1903			
115	EST00099	AONE	EST02115				
124	EST00107	1115	EST02135	1904	EST01019		•
128	EST00252	1112	E3102133	1914 1930 1944	EST01026		
		1110	EST02138	1930	EST01040		
	EST00130	1129	EST02149	1944	EST01050		
164	EST00135	1133	EST02153	1949	EST01054		•
	EST00137	1141	EST02163	. 1962	EST01062		
	EST00296	. 1163	EST02187	1973	EST01071		
179	EST00145	1183	EST02208	1077	FSTR1075		
183	EST00148	1243	EST02272	1982	EST01080		
201	EST00163	1264	EST02293	1001	EST01088		
205	EST00165	1265	EST02294	1003	EST01090		
215	EST00172	. 1266	EST02295	3000	ECT01070		•
230	EST00181		EST02317	1982 1991 1993 2000 2001 2012	EST01097		•
253	EST00199	1308	E3102317	2001	EST01098		•
263	EST00203			2012	EST01106		
		1324	EST02354	2013	EST01107	•	
268	EST00369	1344	EST02374	2024	EST01117		
270	EST00207	1,300	53105700	2013 2024 2043 2051	EST01131		•
271	EST00283	1365	EST02396	2051	EST01138		
273	EST00208	1383	EST02415	2056	EST01142		
	EST00211	1399	EST02433	2058	EST01144		•
281	EST00214	1401	EST02435	2059	EST01145		
. 285	EST00286	1405	EST02439	2064	EST01149		
333	EST00394	1417	EST02452	2000	EST01167		
336	EST00397	1451	EST02487	2056 2058 2059 2064 2090 2094	EST01171		
339	EST00400	1457	EST02493	2116	EST01192		
362	EST00418	1463	EST02500	2117	E0101172		
389	EST00440	1473	EST02510	2117		•	
441	EST00481	1473	EST02516	2128 2131 2134 2144 2145 2150 2151	EST01202		
	EST00493	1514	ESTUZSIO	2131	EST01205		
	EST00509	. 1516	EST02555	2134	EST01208		
	EST00522	1528		2144	EST01216		
		1531	EST02572	2145	EST01217		
504	EST00529	. 1544	EST02586	2150	EST01222		•
	EST00538	1551	EST02593	2155	EST01227		
	EST00540	1558	EST02601	2:101	EST01231		
551	EST01482	1561	EST02604 EST02625 EST02631	21.3	EST01238		
	EST00565	1581	EST02625	2174	EST01242		
559	EST00570	1586	EST02631	2176	EST01244		•
582	EST00592	. 1591	EST02636	2189	EST01255		
602	EST00606	1616	EST02661	2214	EST01272		
606	EST00609	1624	EST02670	2225	EST01278		
608	EST00611	1630	EST02676	2227	EST01279		
621	EST00620	1637	EST00796	2227 2233 2235	EST01284		•
635	EST00629	1639		2235			
	EST00634		EST00808		EST01286		
644	EST00636	1651	EST00810	2236	EST01287	•	
687	EST00671			2255	EST01302		
700		1677		2259	EST01304		
	EST00683		EST00839	2263	EST01307	•	
743	EST00714		EST00849	SEQ ID#	EST#	•	
<i>7</i> 53	EST00721		EST00857	•			
	EST00726		EST00858	2267	EST01756		•
764	EST00729		EST00860	. 2281	EST01321		
808	EST00761	1716	EST00865		EST01322		
823	EST01864	SEQ_ID#	EST#	2300	EST01333		
834	EST00771				EST01335		
886	EST01886		EST00867	2707	EST01335		
919			EST00879	2314			•
	EST01933		EST00887		ESTO1345		•
SEQ ID#			EST00891	2334			
<u> </u>					EST01362	•	
936	ECT01070		EST00903		EST01365		
	EST01939		EST00907	2348	EST01371		
948 945	EST01957		EST00909	2358	EST01379		<u> </u>
965	EST01978	1777	EST00913	2367	EST01388		think.
•							

Table 7: ESTs with Good Pr bability f C ntaining Coding Sequence

SEO ID#	EST#	1041	EST02057		2362	EST01383
		1083	EST02102		2378	EST01397
20	EST00024	1099	EST02118		2399	EST01423
72	EST00071	1105	EST02124		2407	EST02714
82	EST00078	1113	EST02133		2407	E0102/14
88	EST00084	1139	EST02161			
137	EST00272	1146	EST02168			
177	EST00328	1196	EST02100			
193	EST00156	1210	EST02238			
200	EST00162	1233	EST02256			
218	EST00102 EST00175	1233	EST02262 EST02314			
. 228	EST00179	1331	EST02314 EST02361			
247	EST00279	1388	EST02361 EST02421			
264	EST00204	1418	EST02421			
267	EST00204	1439	EST02455			
296	EST00297	1502	EST02540			
371	EST00226	1537				
385		1563	EST02578			
392	EST00436		EST02606			
-	EST00442	1599	EST02644			
414	EST00460	1602	EST02647			
433	EST00474	1693	EST00848			
453	EST00492	1695	EST00850			•
471	EST00505	1729	EST00877			
496	EST00525	1730	EST00878	*	•	
· 524	EST00544	1738	EST00883			
526 529	EST00546	1739	EST00885			
	EST00549	1743	EST00888			
549 557	EST00563	1768	EST00908			
557 578	EST00569	1780	EST00916			
578 ·		1804	EST00938			
596	EST00602	1805	EST00939			
607	EST00610	. 1811	EST00945			,
619 657	EST00619	1819	EST00950			
660	EST00646 EST00649	1826 1830	EST00956			
689	EST00673		EST00959			
695	EST00673	1845	EST00971			
699	EST00679	1848	EST00974			
729	EST00703	1853	EST00977			•
742	EST00703	1967	EST01066 EST01089	•		
747	EST00717	1992 1994	EST01089	•		
755	EST00717					
· 759	EST00725	SEO ID#	EST#			
776	EST00728	1997	EST01094			
778	EST00740	2046	EST01034 EST01134			
782	EST01551		EST01134 EST01177			
829	EST00768	2101 2102	EST01177 EST01178			
835	EST00768	2102	EST01176 EST01181			
836	EST00772	2105				
862	EST01872		EST01182			
881	EST01872 EST01881	2141	EST01213			
SEQ ID#		2184	EST01251			
SEC ID#	EST#	2196	EST01260			
004	E0001004	2203	EST01264			
884	EST01884	2232	EST01283			
924	EST01926	2308	EST01339			
929	EST01932	2345	EST01368			
938	EST01941	2346	EST01369			
971	EST01985	2351	EST01373			
995	EST02009	2354	EST01375	•	•	•
996	EST0201C	2355	EST01376	*		ू इंडर पहार - इंडर पहार
1031	EST02046	2359	EST01380			• •

Table 8: ESTs with Marginal Probability of Containing Coding Sequence

12 EST00274 1267 EST0229 24 EST00027 1301 EST0233 45 EST00364 1397 EST0243 79 EST00076 1448 EST0248 90 EST00302 1480 EST0251 110 EST00096 1493 EST0253 44 EST00120 1499 EST0253 45 EST00121 1503 EST0254 45 EST00121 1503 EST0254 46 EST00121 1503 EST0254 47 EST00121 1503 EST0254 48 EST00184 1548 EST0259 277 EST00212 1562 EST0260 319 EST00381 1572 EST0261 368 EST00423 1575 EST0264 387 EST00425 1595 EST0264 387 EST00425 1595 EST0264 402 EST00451 1610 EST0265 415 EST00464 1627 EST0266 418 EST00464 1627 EST0267 503 EST00580 1683 EST0084 532 EST00528 1683 EST0084 532 EST00551 1751 EST0089 570 EST00580 1764 EST0269 573 EST00580 1764 EST0090 570 EST00580 1764 EST0090 570 EST00580 1764 EST0090 617 EST00617 1877 EST0099 618 EST00615 1900 EST0101 738 EST00711 1954 EST0097 726 EST00700 1939 EST0165 727 EST00710 1940 EST0104 738 EST00715 SEO ID# EST0107 745 EST00777 2041 EST0103 852 EST00756 2031 EST0103 854 EST00777 2041 EST0103 854 EST00777 2041 EST0103 855 EST00784 2008 EST01103 854 EST00777 2041 EST0113 855 EST00784 2008 EST01103 854 EST00777 2041 EST0113 855 EST00784 2060 EST01103 856 EST01990 2008 EST01103 857 EST01907 2100 EST0117 852 EST00778 2044 EST0113 854 EST00718 2044 EST0113 855 EST01990 2050 EST01129 935 EST01938 2153 EST0129 935 EST01991 2250 EST0129 938 EST01991 2250 EST0129							
11 EST00018 1224 EST0225 12 EST00274 1267 EST0229 24 EST0027 1301 EST0223 45 EST00364 1397 EST0243 79 EST00076 1448 EST0248 90 EST00302 1480 EST0253 110 EST00096 1493 EST0253 144 EST00120 1499 EST0253 145 EST00121 1503 EST0254 1502 EST00155 1527 EST0256 222 EST00177 1536 EST0257 234 EST00184 1548 EST0259 277 EST00212 1562 EST0260 319 EST00381 1572 EST0261 368 EST00423 1575 EST0261 368 EST00423 1575 EST0264 387 EST00425 1595 EST0264 415 EST00461 1621 EST0266 415 EST00461 1621 EST0266 418 EST00461 1621 EST0266 418 EST00464 1627 EST0267 503 EST00528 1631 EST0267 504 EST00539 1683 EST0084 522 EST00551 1751 EST0089 532 EST00580 1764 EST0269 540 EST00580 1764 EST0269 570 EST00580 1764 EST0269 570 EST00586 1793 EST0092 613 EST00615 1847 EST0097 626 EST00665 1900 EST0101 738 EST00615 1847 EST0097 626 EST00700 1939 EST0101 738 EST00711 1954 EST0104 738 EST00711 1954 EST0104 738 EST00711 1954 EST0101 745 EST00749 2008 EST0113 852 EST00756 2031 EST012 955 EST00782 2044 EST0113 854 EST00715 SEO ID# EST# 550 ID# EST# 2248 EST0129 968 EST01999 2250 EST0129 968 EST01999 2250 EST0129 968 EST01999 2250 EST0129	SEQ	ID#	EST#		1222	EST02251	
11 EST00018							
12 EST00274 1267 EST0229 24 EST00027 1301 EST0233 45 EST00364 1397 EST0248 79 EST00076 1448 EST0248 90 EST00302 1480 EST0251 110 EST00096 1493 EST0253 44 EST00120 1499 EST0253 45 EST00121 1503 EST0254 45 EST00121 1503 EST0254 46 EST00121 1503 EST0255 222 EST00177 1536 EST0257 234 EST00184 1548 EST0259 277 EST00212 1562 EST0260 319 EST00381 1572 EST0261 368 EST00423 1575 EST0264 387 EST00425 1595 EST0264 387 EST00425 1595 EST0264 402 EST00464 1608 EST0265 418 EST00464 1627 EST0266 418 EST00464 1627 EST0267 503 EST00528 1631 EST0267 503 EST00528 1631 EST0267 504 EST0059 1683 EST0084 532 EST0051 1751 EST0084 532 EST00551 1751 EST0089 570 EST00580 1764 EST0269 570 EST00580 1764 EST0269 613 EST00586 1793 EST0090 570 EST00586 1793 EST0090 617 EST00617 1877 EST0091 626 EST00617 1877 EST0097 626 EST00622 1897 EST0101 738 EST00615 1900 EST0101 738 EST00711 1940 EST0104 738 EST00710 1940 EST0104 738 EST00711 1954 EST0105 727 EST00700 1939 EST0165 728 EST00777 2041 EST0103 852 EST00756 2031 EST0103 854 EST00777 2041 EST0113 852 EST00777 2041 EST0113 852 EST00784 2060 EST01103 854 EST00777 2041 EST0113 855 EST00784 2060 EST01103 856 EST0099 2008 EST01129 935 EST01938 2153 EST0129 935 EST01938 2153 EST0129 938 EST01999 92250 EST0129 938 EST01999 9250 EST0129		11	EST00018			EST02257	
24 EST00027 1301 EST0233 45 EST00364 1397 EST0243 79 EST00076 1448 EST0243 90 EST00302 1480 EST0251 110 EST00096 1493 EST0253 1446 EST00120 1499 EST0253 45 EST00121 1503 EST0254 1503 EST0254 1503 EST0255 1527 EST0256 1527 EST00256 1527 EST00256 1527 EST00256 1527 EST00256 1527 EST00261 1503 EST0257 1536 EST0257 1536 EST0257 1536 EST0261 1539 EST00423 1575 EST0261 1536 EST0265 1527 EST00425 1595 EST0263 1595 EST0264 155 EST00425 1595 EST0265 1415 EST00451 1610 EST0265 1415 EST00461 1621 EST0266 1418 EST00464 1627 EST0267 1629 EST0267 1629 EST00267 1629 EST00267 1629 EST00267 1629 EST00267 1629 EST00267 1629 EST00267 1629 EST0084 1532 EST00539 1683 EST0084 1532 EST00539 1683 EST0084 1532 EST00557 1756 EST0089 1540 EST00557 1756 EST0089 1570 EST00580 1764 EST0269 1770 EST0090 1764 EST0097 1766 EST0097 1766 EST0097 1766 EST00615 1847 EST0097 1627 EST00617 1877 EST0097 1627 EST00617 1877 EST0097 1628 EST00101 1939 EST0105 1755 EST0065 1900 EST0101 1755 EST00065 1900 EST0101 1755 EST00072 1940 EST0101 1954 EST0105 1755 EST00709 1939 EST0105 1755 EST00709 1939 EST0105 1755 EST00709 1939 EST0105 1755 EST00709 1930 EST0		12	EST00274				
45 EST00364 1397 EST0243 79 EST00076 1448 EST0248 90 EST00302 1480 EST0253 110 EST00096 1493 EST0253 144 EST00120 1499 EST0253 145 EST00121 1503 EST0254 192 EST00155 1527 EST0256 222 EST00177 1536 EST0257 234 EST00184 1548 EST0259 277 EST00212 1562 EST0261 368 EST00423 1575 EST0261 370 EST00425 1595 EST0264 387 EST00425 1595 EST0264 415 EST00461 1621 EST0265 415 EST00461 1621 EST0265 415 EST00464 1627 EST0267 426 EST00470 1629 EST0267 503 EST00528 1631 EST0267 517 EST00539 1683 EST00267 520 EST00557 1756 EST0090 570 EST00586 1793 EST0099 570 EST00586 1793 EST0099 613 EST00586 1793 EST0099 614 EST00615 1847 EST0099 615 EST00615 1847 EST0099 616 EST00617 1877 EST0099 617 EST00615 1847 EST0099 618 EST00615 1900 EST0101 726 EST00617 1877 EST0099 618 EST00700 1939 EST0105 726 EST00700 1939 EST0105 727 EST00701 1940 EST0104 738 EST00711 1954 EST0104 745 EST00746 1990 EST0101 745 EST00746 1990 EST0101 745 EST00746 1990 EST0101 745 EST00746 1990 EST0104 745 EST00777 2041 EST0105 752 EST00749 2008 EST01102 845 EST00777 2041 EST0112 852 EST00784 2060 EST01102 968 EST01990 2008 EST01126 968 EST01991 2016 EST01269 968 EST01991 2250 EST01299 968 EST01999 9250 EST01299 968 EST01999 9250 EST01299		24	EST00027			EST02331	
79 EST00076 90 EST00302 1480 EST0251 110 EST00096 1493 EST0253 144 EST00120 1499 EST0253 145 EST00121 1503 EST0254 192 EST00155 1527 EST0256 222 EST00177 1536 EST0257 234 EST00184 1548 EST0259 277 EST00212 1562 EST0261 368 EST00423 370 EST00425 387 EST00425 1595 EST0264 415 EST00461 1610 EST0265 415 EST00461 4162 EST00464 416 EST00464 418 EST00464 41627 EST0267 503 EST00528 1631 EST0267 503 EST00528 1631 EST0088 522 EST00539 1663 EST0088 532 EST00551 1751 EST0089 570 EST00580 1764 EST0090 570 EST00586 1793 EST0091 576 EST00586 1793 EST0092 681 EST00617 1877 EST0091 617 EST00617 1877 EST0091 618 EST00615 619 EST0065 726 EST00665 1900 EST0101 738 EST00711 1954 EST0101 745 EST00715 752 EST00720 791 EST00746 1990 EST0108 803 EST00756 2031 EST0112 854 EST00784 2060 EST0112 855 EST01999 988 EST01202 2266 EST01299 988 EST01202 2266 EST01299 988 EST01202 2266 EST01299 988 EST01202 2266 EST01290 988 EST01202 2266 EST01230	•	45	EST00364	**		EST02431	
90 EST00302 1480 EST0251 110 EST00096 1493 EST0253 444 EST00120 1499 EST0253 45 EST00121 1503 EST0254 45 EST00155 1527 EST0256 222 EST00177 1536 EST0257 234 EST00184 1548 EST0259 277 EST00212 1562 EST0260 319 EST00381 1572 EST0261 368 EST00423 1575 EST0261 370 EST00425 1595 EST0264 402 EST00451 1610 EST0265 415 EST00461 1621 EST0266 418 EST00464 1627 EST0267 426 EST00470 1629 EST0267 503 EST00528 1631 EST0267 517 EST00539 1683 EST00267 517 EST00539 1683 EST0084 532 EST00551 1751 EST0089 532 EST00580 1764 EST0269 573 EST00580 1764 EST0099 570 EST00580 1764 EST0099 570 EST00580 1764 EST0099 617 EST00617 1877 EST0099 618 EST00665 1990 EST0101 626 EST00665 1990 EST0101 738 EST0071 1940 EST0104 738 EST0071 1940 EST0104 738 EST0071 1990 EST0104 745 EST00756 1990 EST0105 745 EST00756 1990 EST0108 803 EST00756 2031 EST0129 804 EST00764 1990 EST0104 854 EST00777 2041 EST0113 855 EST00784 2060 EST01105 855 EST00784 2060 EST01107 912 EST00784 2060 EST01107 912 EST00784 2060 EST01126 854 EST00777 2041 EST01136 855 EST00784 2060 EST01126 854 EST00784 2060 EST01126 935 EST01991 2016 EST01296 935 EST01991 2250 EST01296 988 EST01299 2250 EST01299 988 EST02002 2266 EST01330		79	EST00076			EST02484	
110 EST00096	•	90	EST00302				
### ### ### ### ### ### ### ### ### ##		110	EST00096				
### ### ### ### ### ### ### ### ### ##		.44	EST00120			EST02537	
192 EST00155 1527 EST0256		45	EST00121	•		EST02541	
222 EST00177 234 EST00184 277 EST00212 319 EST00381 368 EST00423 370 EST00425 387 EST00425 402 EST00451 415 EST00461 416 EST00464 418 EST00464 418 EST00464 418 EST00464 418 EST00466 418 EST00528 517 EST00528 517 EST00539 522 EST00539 532 EST00551 532 EST00551 533 EST00557 540 EST00580 573 EST00580 574 EST00580 575 EST00580 576 EST00586 577 EST00586 578 EST00665 579 EST00665 726 EST00665 726 EST00700 727 EST00710 728 EST00710 729 EST00720 791 EST00746 795 EST00749 803 EST00756 803 EST00756 803 EST00777 804 EST00777 805 EST00784 805 EST00794 806 EST00794 807 EST0097 807 EST00794 808 EST00794 809 EST01133 856 EST00794 907 EST01912 918 EST01269 988 EST01999 988 EST01299		-92	EST00155				
234 EST00184 1548 EST0259 277 EST00212 1562 EST0260 319 EST00381 1572 EST0261 368 EST00423 1575 EST0261 370 EST00425 1595 EST0264 387 EST00438 1608 EST0265 402 EST00451 1610 EST0265 415 EST00461 1621 EST0266 418 EST00464 1627 EST0267 426 EST00470 1629 EST0267 503 EST00528 1631 EST0267 517 EST00539 1683 EST0084 522 EST00551 1751 EST0089 522 EST00551 1751 EST0089 532 EST00557 1756 EST0090 570 EST00580 1764 EST0269 573 EST00580 1764 EST0269 573 EST00586 1793 EST0092 613 EST00615 1847 EST0097 617 EST00617 1877 EST0097 626 EST00665 1900 EST0101 726 EST00700 1939 EST0101 727 EST00711 1954 EST0105 728 EST00711 1954 EST0105 729 EST00746 1990 EST0101 738 EST00746 1990 EST0104 738 EST00756 2031 EST0129 791 EST00746 1990 EST0113 854 EST00777 2041 EST0113 854 EST00777 2041 EST0113 854 EST00784 2060 EST0110 803 EST00756 2031 EST0112 935 EST01938 2153 EST0129 935 EST01938 2153 EST0129 938 EST01999 2250 EST0129 988 EST01999 2250 EST0129 988 EST01999 2250 EST0129	•	222	EST00177		1536		
277 EST00212 1562 EST0260 319 EST00381 1572 EST0261 368 EST00423 1575 EST0261 370 EST00425 1595 EST0264 387 EST00451 1608 EST0265 402 EST00451 1610 EST0265 415 EST00461 1621 EST0266 418 EST00464 1627 EST0267 503 EST00528 1631 EST0267 517 EST00539 1683 EST0084 522 EST00539 1683 EST0084 532 EST00551 1751 EST0089 540 EST00557 1756 EST0090 570 EST00580 1764 EST0269 573 EST00586 1793 EST0092 613 EST00586 1793 EST0092 614 EST00615 1847 EST0099 615 EST00665 1900 EST0101 726 EST00665 1900 EST0101 726 EST00700 1939 EST0165 727 EST00701 1940 EST0104 738 EST00711 1954 EST0105 745 EST00749 2008 EST0104 755 EST00749 2008 EST0110 803 EST00756 2031 EST0112 845 EST00777 2041 EST0113 852 EST00784 2060 EST0110 803 EST00756 2031 EST0112 854 EST00777 2041 EST0113 855 EST00784 2060 EST0110 807 EST01997 2100 EST0117 912 EST01912 2136 EST0122 935 EST01938 2153 EST0122 935 EST01938 2153 EST0122 968 EST01999 2250 EST0129 988 EST01999 2250 EST0129		234	EST00184			EST02590	
319 EST00381 1572 EST0261 368 EST00423 1575 EST0261 370 EST00425 1595 EST0264 387 EST00438 1608 EST0265 402 EST00451 1610 EST0265 415 EST00461 1621 EST0266 418 EST00464 1627 EST0267 426 EST00470 1629 EST0267 503 EST00528 1631 EST0267 517 EST00539 1683 EST0084 522 EST00543 1692 EST0084 532 EST00551 1751 EST0089 540 EST00557 1756 EST0090 570 EST00580 1764 EST0269 573 EST00583 1770 EST0091 576 EST00586 1793 EST0092 613 EST00586 1793 EST0092 614 EST00615 1847 EST0097 617 EST00617 1877 EST0099 618 EST00665 1900 EST0101 726 EST00700 1939 EST0165 727 EST00701 1940 EST0101 738 EST00711 1954 EST0105 745 EST00720 791 EST00746 1990 EST0108 795 EST00749 2008 EST01102 803 EST00756 2031 EST0112 845 EST00777 2041 EST0113 852 EST00784 2060 EST01102 854 EST00784 2060 EST0113 855 EST01999 208 EST0122 935 EST01938 2153 EST0122 935 EST01938 2153 EST0122 936 EST01999 92250 EST0129 988 EST01999 2250 EST0129		277	EST00212			EST02605	
368 EST00423 1575 EST0261 370 EST00425 1595 EST0264 387 EST00438 1608 EST0265 402 EST00451 1610 EST0265 415 EST00461 1621 EST0266 418 EST00464 1627 EST0267 503 EST00528 1631 EST0267 503 EST00539 1683 EST0084 522 EST00543 1692 EST0084 532 EST00551 1751 EST0089 540 EST00557 1756 EST0090 570 EST00580 1764 EST0269 573 EST00583 1770 EST0091 576 EST00586 1793 EST0092 613 EST00615 1847 EST0097 617 EST00617 1877 EST0099 626 EST00622 1897 EST0101 681 EST00665 1900 EST0101 726 EST00700 1939 EST0105 727 EST00701 1940 EST0104 738 EST00711 1954 EST01056 727 EST00702 1930 EST0105 728 EST00715 SEO ID# EST# 752 EST00749 2008 EST0110 803 EST00756 2031 EST0112 845 EST00777 2041 EST01136 845 EST00777 2041 EST01136 845 EST00777 2041 EST01136 852 EST00782 2044 EST01136 854 EST00784 2060 EST01146 907 EST01907 2100 EST01176 912 EST01912 2136 EST0129 935 EST01938 2153 EST0129 948 EST01999 2250 EST0129 948 EST01999 2250 EST0129 948 EST01999 2250 EST0129		319	EST00381			EST02615	
370 EST00425 1595 EST0264 387 EST00438 1608 EST0265 402 EST00451 1610 EST0265 415 EST00461 1621 EST0266 418 EST00464 1627 EST0267 426 EST00470 1629 EST0267 503 EST00528 1631 EST0267 517 EST00539 1683 EST0084 532 EST00551 1751 EST0089 540 EST00557 1756 EST0090 570 EST00580 1764 EST0269 573 EST00583 1770 EST0091 576 EST00586 1793 EST0092 613 EST00586 1793 EST0092 614 EST00615 1847 EST0097 617 EST00617 1877 EST0099 626 EST00622 1897 EST0101 681 EST00665 1900 EST0101 726 EST00700 1939 EST0165 727 EST00710 1940 EST0104 738 EST00711 1954 EST01056 727 EST00740 1990 EST0108 728 EST00746 1990 EST0108 795 EST00746 1990 EST0108 795 EST00749 2008 EST0110 803 EST00756 2031 EST0112 845 EST00777 2041 EST0113 852 EST00782 2044 EST0113 854 EST00784 2060 EST01146 907 EST01907 2100 EST01146 907 EST01907 2100 EST01146 907 EST01912 2136 EST0122 935 EST01938 2153 EST0122 968 EST01981 2248 EST0129 988 EST02002 2266 EST01310		368	EST00423		_	EST02618	
387 EST00438 1608 EST0265		370	EST00425			EST02640	
402 EST00451 1610 EST0265 415 EST00461 1621 EST0266 418 EST00464 1627 EST0267 426 EST00470 1629 EST0267 503 EST00528 1631 EST0267 517 EST00539 1683 EST0084 522 EST00543 1692 EST0084 532 EST00551 1751 EST0089 540 EST00557 1756 EST0090 570 EST00580 1764 EST0269 573 EST00583 1770 EST0091 576 EST00586 1793 EST0092 613 EST00615 1847 EST0097 626 EST00617 1877 EST0099 626 EST00622 1897 EST0101 681 EST00665 1990 EST0101 726 EST00700 1939 EST0105 727 EST00701 1940 EST0104 738 EST00715 SEO ID# EST# 752 EST00720 791 EST00746 1990 EST0105 803 EST00756 2031 EST0112 845 EST00777 2041 EST0112 852 EST00784 2060 EST0114 907 EST01907 2100 EST0112 912 EST01912 2136 EST0121 935 EST01938 2153 EST0122 968 EST01981 2248 EST0129 988 EST01999 2250 EST0129 988 EST01999 2250 EST0129 988 EST01999 2250 EST0129		387	EST00438			EST02653	
415 EST00461 1621 EST0266 418 EST00464 1627 EST0267 426 EST00470 1629 EST0267 503 EST00528 1631 EST0267 517 EST00539 1683 EST0084 522 EST00543 1692 EST0084 532 EST00551 1751 EST0089 540 EST00557 1756 EST0090 570 EST00580 1764 EST0269 573 EST00583 1770 EST0091 576 EST00586 1793 EST0092 613 EST00615 1847 EST0097 617 EST00617 1877 EST0099 626 EST00617 1877 EST0097 626 EST00665 1900 EST0101 726 EST00700 1939 EST0165 727 EST00701 1940 EST0104 738 EST00711 1954 EST0105 745 EST00715 SEQ ID# EST# 752 EST00749 2008 EST01105 803 EST00756 2031 EST0112 845 EST00777 2041 EST0113 852 EST00784 2060 EST0114 907 EST01907 2100 EST0113 854 EST00784 2060 EST0113 855 EST01912 2136 EST0122 935 EST01938 2153 EST0122 968 EST01981 2248 EST0129 968 EST01999 2250 EST0129 968 EST01999 2250 EST0129 968 EST01999 2250 EST0129 968 EST01999 2250 EST0129		402	EST00451			EST02655	
418 EST00464 426 EST00470 503 EST00528 503 EST00528 517 EST00539 522 EST00543 532 EST00551 540 EST00557 570 EST00580 570 EST00580 571 EST00583 572 EST00583 573 EST00583 574 EST00586 573 EST00586 574 EST00615 613 EST00615 614 EST00617 626 EST00622 618 EST00665 727 EST00700 726 EST00701 727 EST00701 728 EST00711 729 EST00720 791 EST00746 795 EST00746 795 EST00746 795 EST00777 852 EST00777 854 EST00784 803 EST00756 804 EST00764 907 EST01907 912 EST01912 935 EST01918 988 EST01981 988 EST01299 988 EST01999 988 EST01299		415	EST00461			EST02667	
426 EST00470 1629 EST0267 503 EST00528 1631 EST0267 517 EST00539 1683 EST0084 522 EST00543 1692 EST0084 532 EST00551 1751 EST0089 540 EST00557 1756 EST0089 570 EST00580 1764 EST0269 573 EST00583 1770 EST0091 576 EST00586 1793 EST0092 613 EST00615 1847 EST0097 617 EST00617 1877 EST0097 626 EST00622 1897 EST0101 681 EST00700 1939 EST0165 727 EST00701 1940 EST0105 726 EST00711 1954 EST0105 727 EST00720 1990 EST0108 791 EST00720 1990 EST0108 795 EST00782 2031 EST0112 845 EST00782 2044 EST0113 854 E		418	EST00464	***	1627	EST02674	
517 EST00539 1683 EST0084 522 EST00543 1692 EST0084 532 EST00551 1751 EST0089 540 EST00557 1756 EST0090 570 EST00580 1764 EST0269 573 EST00583 1770 EST0091 576 EST00586 1793 EST0092 613 EST00615 1847 EST0097 617 EST00617 1877 EST0099 626 EST00622 1897 EST0101 681 EST00655 1900 EST0101 726 EST00700 1939 EST0165 727 EST00701 1940 EST0105 727 EST00711 1954 EST0105 738 EST00715 SEO ID# EST# 752 EST00720 1990 EST0108 795 EST00749 2008 EST0112 803 EST00777 2041 EST0113 854 EST00784 2060 EST01146 907 EST01991		426	EST00470		1629	EST02677	•
522 EST00543 1692 EST0084 532 EST00551 1751 EST0089 540 EST00557 1756 EST0090 570 EST00580 1764 EST0269 573 EST00583 1770 EST0091 576 EST00586 1793 EST0092 613 EST00615 1847 EST0097 617 EST00617 1877 EST0099 626 EST00622 1897 EST0101 681 EST00665 1900 EST0101 726 EST00700 1939 EST0165 727 EST00701 1940 EST0105 727 EST00715 SEQ ID# EST# 752 EST00720 1994 EST0108 795 EST00746 1990 EST0108 795 EST00749 2008 EST0112 854 EST00777 2041 EST0113 854 EST01990 2044 EST0113 907 EST01902 206 EST01176 912 E		503	EST00528		1631	EST02678	
532 EST00551 1751 EST0089 540 EST00557 1756 EST0090 570 EST00580 1764 EST0269 573 EST00586 1793 EST0091 576 EST00586 1793 EST0092 613 EST00615 1847 EST0097 626 EST00622 1897 EST0101 681 EST00665 1900 EST0165 726 EST00700 1939 EST0165 727 EST00701 1940 EST0104 738 EST00711 1954 EST0105 745 EST00715 SEO ID# EST# 752 EST00720 1990 EST0108 791 EST00746 1990 EST0108 795 EST00749 2008 EST0112 845 EST00777 2041 EST0113 852 EST01907 200 EST01146 907 EST01907 2100 EST01176 912 EST01938 2153 EST01226 SEO ID#		517	EST00539		1683	EST00840	
532 EST00551 1751 EST0089 540 EST00557 1756 EST0090 570 EST00580 1764 EST0269 573 EST00583 1770 EST0091 576 EST00586 1793 EST0092 613 EST00615 1847 EST0092 617 EST00617 1877 EST0099 626 EST00665 1900 EST0101 726 EST00700 1939 EST0165 727 EST00701 1940 EST0104 738 EST00711 1954 EST0105 745 EST00715 SEO ID# EST# 752 EST00720 1990 EST0108 791 EST00746 1990 EST0108 795 EST00749 2008 EST0112 845 EST00777 2041 EST0113 854 EST00784 2060 EST01146 907 EST01991 2136 EST0122 935 EST01938 2153 EST0122 968			EST00543		1692	EST00847	
570 EST00580 1764 EST0269 573 EST00583 1770 EST0091 576 EST00586 1793 EST0092 613 EST00615 1847 EST0097 617 EST00617 1877 EST0099 626 EST00622 1897 EST0101 681 EST00665 1900 EST0101 726 EST00700 1939 EST0165 727 EST00701 1940 EST0104 738 EST00711 1954 EST0105 745 EST00715 SEO ID# EST# 752 EST00720 1990 EST0108 791 EST00746 1990 EST0108 795 EST00749 2008 EST0112 803 EST00756 2031 EST0112 845 EST00777 2041 EST0113 854 EST00784 2060 EST01146 907 EST01907 2100 EST0122 935 EST01938 2153 EST0122 968					1751	EST00895	
573 EST00583 1770 EST0091 576 EST00586 1793 EST0092 613 EST00615 1847 EST0097 617 EST00617 1877 EST0099 626 EST00622 1897 EST0101 681 EST00665 1900 EST0101 726 EST00700 1939 EST0165 727 EST00701 1940 EST0104 738 EST00711 1954 EST0105 745 EST00715 SEO ID# EST# 752 EST00720 791 EST00746 1990 EST0108 795 EST00749 2008 EST0110 803 EST00756 2031 EST0112 845 EST00777 2041 EST0113 852 EST00782 2044 EST0113 854 EST00784 2060 EST0114 907 EST01907 2100 EST0117 912 EST01912 2136 EST0121 935 EST01938 2153 EST0122 935 EST01938 2153 EST0122 968 EST01981 2248 EST0129 988 EST02002 2266 EST01310					1756	EST00900	
576 EST00586 1793 EST0092 613 EST00615 1847 EST0097 617 EST00617 1877 EST0099 626 EST00622 1897 EST0101 681 EST00765 1900 EST0101 726 EST00700 1939 EST0165 727 EST00701 1940 EST0104 738 EST00715 SEO ID# EST0105 745 EST00720 EST01# EST# 752 EST00720 EST0108 EST0108 791 EST00746 1990 EST0108 795 EST00749 2008 EST0112 803 EST00756 2031 EST0112 845 EST00777 2041 EST0113 854 EST01982 2044 EST0113 854 EST01907 2100 EST01176 912 EST01912 2136 EST01226 935 EST01938 2153 EST01226 968 EST01981 2248 EST01296 985					1764	EST02690	
613 EST00615 1847 EST0097 617 EST00617 1877 EST0099 626 EST00622 1897 EST0101 681 EST00665 1900 EST0101 726 EST00700 1939 EST0165 727 EST00701 1940 EST0104 738 EST00711 1954 EST0105 745 EST00715 SEO ID# EST# 752 EST00720 791 EST00746 1990 EST0108 795 EST00749 2008 EST0110 803 EST00756 2031 EST0112 845 EST00777 2041 EST0113 852 EST00782 2044 EST0113 854 EST00784 2060 EST0114 907 EST01907 2100 EST0117 912 EST01907 2100 EST0117 912 EST01912 2136 EST01212 935 EST01938 2153 EST0122 935 EST01938 2153 EST0122 935 EST01981 2248 EST0129 968 EST01999 2250 EST0129 988 EST02002 2266 EST01310					1770	EST00910	
617 EST00617 1877 EST0099 626 EST00622 1897 EST01013 681 EST00665 1900 EST01013 726 EST00700 1939 EST0165 727 EST00701 1940 EST0104 738 EST00711 1954 EST0105 745 EST00715 SEO ID# EST# 752 EST00720 791 EST00746 1990 EST0108 795 EST00749 2008 EST01103 803 EST00756 2031 EST0112 845 EST00777 2041 EST0113 852 EST00782 2044 EST0113 854 EST00784 2060 EST0114 907 EST01907 2100 EST01176 912 EST01907 2100 EST01176 912 EST01912 2136 EST01219 935 EST01938 2153 EST0122 935 EST01938 2153 EST0122 968 EST01981 2248 EST0129 968 EST01999 2250 EST0129 988 EST02002 2266 EST01310	•					EST00929	
626 EST00622 1897 EST01013 681 EST00665 1900 EST01013 726 EST00700 1939 EST0165 727 EST00701 1940 EST01046 738 EST00711 1954 EST01056 745 EST00715 SEO ID# EST# 752 EST00720 791 EST00746 1990 EST0108 795 EST00749 2008 EST01103 803 EST00756 2031 EST01123 845 EST00777 2041 EST01133 852 EST00782 2044 EST01133 854 EST00784 2060 EST01146 907 EST01907 2100 EST01176 912 EST01907 2100 EST01176 912 EST01912 2136 EST01223 935 EST01938 2153 EST01223 SEO ID# EST# 2204 EST01263 2212 EST01276 968 EST01981 2248 EST01293 985 EST01999 2250 EST01293 988 EST02002 2266 EST01310					1847	EST00973	
681 EST00665 1900 EST01019 726 EST00700 1939 EST01659 727 EST00701 1940 EST01040 738 EST00711 1954 EST01050 745 EST00715 SEO ID# EST# 752 EST00720 791 EST00746 1990 EST01080 795 EST00749 2008 EST01101 803 EST00756 2031 EST01120 845 EST00777 2041 EST01130 852 EST00782 2044 EST01130 854 EST00784 2060 EST01140 907 EST01907 2100 EST01170 912 EST01912 2136 EST01210 935 EST01938 2153 EST01220 948 EST01981 2248 EST01290 988 EST02002 2266 EST01310					1877	EST00998	
726 EST00700 1939 EST01655 727 EST00701 1940 EST01046 738 EST00711 1954 EST01056 745 EST00715 SEO ID# EST# 752 EST00720 791 EST00746 1990 EST0108 795 EST00749 2008 EST01105 803 EST00756 2031 EST01125 845 EST00777 2041 EST01136 852 EST00782 2044 EST01136 854 EST00784 2060 EST01146 907 EST01907 2100 EST01176 912 EST01912 2136 EST01216 935 EST01938 2153 EST01225 935 EST01938 2153 EST01225 968 EST01981 2248 EST01295 988 EST02002 2266 EST01316					1897	EST01012	
727 EST00701 1940 EST01040 738 EST00711 1954 EST01050 745 EST00715 SEQ ID# EST# 752 EST00720 791 EST00746 1990 EST0108 795 EST00749 2008 EST01103 803 EST00756 2031 EST01123 845 EST00777 2041 EST01130 852 EST00782 2044 EST01133 854 EST00784 2060 EST01146 907 EST01907 2100 EST01176 912 EST01912 2136 EST01212 935 EST01938 2153 EST01223 SEO ID# EST# 2204 EST01223 968 EST01981 2248 EST01293 988 EST02002 2266 EST01310						EST01015	
738 EST00711 1954 EST01056 745 EST00715 SEQ ID# EST# 752 EST00720 791 EST00746 1990 EST0108 795 EST00749 2008 EST01103 803 EST00756 2031 EST01123 845 EST00777 2041 EST01133 852 EST00782 2044 EST01133 854 EST00784 2060 EST01146 907 EST01907 2100 EST01176 912 EST01912 2136 EST01212 935 EST01938 2153 EST01223 SEQ ID# EST# 2204 EST01223 968 EST01981 2248 EST01293 988 EST02002 2266 EST01310						EST01655	
745 EST00715 752 EST00720 791 EST00746 1990 EST0108 795 EST00749 803 EST00756 845 EST00777 852 EST00777 2041 EST0112 854 EST00782 854 EST00784 907 EST01907 912 EST01907 912 EST01912 935 EST01912 935 EST01938 2153 EST0122 968 EST01981 968 EST01981 985 EST01999 988 EST02002 2266 EST01310	٠.					EST01046	
752 EST00720 791 EST00746 1990 EST0108 795 EST00749 2008 EST01103 803 EST00756 2031 EST01123 845 EST00777 2041 EST01133 852 EST00782 2044 EST01133 854 EST00784 2060 EST01146 907 EST01907 2100 EST01176 912 EST01912 2136 EST01213 935 EST01938 2153 EST01223 SEO ID# EST# 2204 EST01223 968 EST01981 2248 EST01293 988 EST02002 2266 EST01310						EST01058	
791 EST00746 1990 EST0108 795 EST00749 2008 EST0110 803 EST00756 2031 EST0112 845 EST00777 2041 EST0113 852 EST00782 2044 EST0113 854 EST00784 2060 EST01146 907 EST01907 2100 EST01176 912 EST01912 2136 EST01210 935 EST01938 2153 EST01225 SEO ID# EST# 2204 EST01265 968 EST01981 2248 EST0129 988 EST02002 2266 EST01310					SEO ID#	EST#	
795 EST00749 2008 EST01101 803 EST00756 2031 EST01121 845 EST00777 2041 EST01131 852 EST00782 2044 EST01133 854 EST00784 2060 EST01146 907 EST01907 2100 EST01176 912 EST01912 2136 EST01210 935 EST01938 2153 EST01220 935 EST01938 2153 EST01220 948 EST01981 2248 EST01290 988 EST01999 2250 EST01290 988 EST02002 2266 EST01310							
803 EST00756 2031 EST01123 845 EST00777 2041 EST01133 852 EST00782 2044 EST01133 854 EST00784 2060 EST01146 907 EST01907 2100 EST01176 912 EST01912 2136 EST01213 935 EST01938 2153 EST01223 SEO ID# EST# 2204 EST0126 968 EST01981 2248 EST01293 985 EST01999 2250 EST01293 988 EST02002 2266 EST01310							
845 EST00777 2041 EST01130 852 EST00782 2044 EST01133 854 EST00784 2060 EST01146 907 EST01907 2100 EST01176 912 EST01912 2136 EST01210 935 EST01938 2153 EST01220 SEO ID# EST# 2204 EST01260 968 EST01981 2248 EST01297 985 EST01999 2250 EST01299 988 EST02002 2266 EST01310			-				
852 EST00782 2044 EST01132 854 EST00784 2060 EST01146 907 EST01907 2100 EST01176 912 EST01912 2136 EST01210 935 EST01938 2153 EST01220 SEO ID# EST# 2204 EST01260 2212 EST01270 968 EST01981 2248 EST01290 985 EST01999 2250 EST01290 988 EST02002 2266 EST01310							
854 EST00784 2060 EST01146 907 EST01907 2100 EST01176 912 EST01912 2136 EST01210 935 EST01938 2153 EST01220 SEO ID# EST# 2204 EST01260 2212 EST01270 968 EST01981 2248 EST01290 985 EST01999 2250 EST01290 988 EST02002 2266 EST01310				٠.,			
907 EST01907 2100 EST01176 912 EST01912 2136 EST01210 935 EST01938 2153 EST01222 SEO ID# EST# 2204 EST01262 968 EST01981 2248 EST01292 985 EST01999 2250 EST01292 988 EST02002 2266 EST01310					_		
912 EST01912 2136 EST01210 935 EST01938 2153 EST01220 SEO ID# EST# 2204 EST01260 2212 EST01270 968 EST01981 2248 EST01290 985 EST01999 2250 EST01290 988 EST02002 2266 EST01310	•			,			
935 EST01938 2153 EST01229 SEO ID# EST# 2204 EST01269 2212 EST01270 968 EST01981 2248 EST01299 985 EST01999 2250 EST01299 988 EST02002 2266 EST01310						EST01176	
SEO ID# EST# 2204 EST01269 2212 EST01270 968 EST01981 2248 EST01299 985 EST01999 2250 EST01299 988 EST02002 2266 EST01310							
968 EST01981 2248 EST0129 985 EST01999 2250 EST0129 988 EST02002 2266 EST01310	SEO						
968 EST01981 2248 EST0129 985 EST01999 2250 EST0129 988 EST02002 2266 EST01310	200	II	<u> </u>				
985 EST01999 2250 EST01299 988 EST02002 2266 EST01310		968	ESTOLOGI				
988 EST02002 2266 EST01310							
				,			•
	1	•				EST01310 EST01340	
2505 25101540						EST01340 EST01370	
				•			
2000 2010120						EST01406 EST01422	
						EST01422 EST01427	
1205 EST02233				•	2403	~01U142/	

Table 9: ESTs with Poor Coding Probability

SEQ ID:	EST#	103	EST00317	204	EST00235	309	EST00174	404	EST00453
		104	EST00354	206		315			
	I EST00007	105		207				405	
		107		209		316		406	
3						317		407	EST00456
		109		210		318		408	EST00457
4		111		211		320	EST00382	409	EST01444
5		112	EST00318	212	EST00169	321	EST00383	410	
	S EST00013	113	EST00097	213	EST00170	322		411	EST00459
8	B EST00234	116	EST00100	214		323			
10		117		216				412	EST01445
14		118				325	EST00386	416	EST00462
				219		326		417	EST00463
. 16		119		220		327	EST00388	419	EST00465
17		120		221	EST00359	328	EST00389	420	EST00466
18	EST00373	121	EST00104	224	EST00356	329	EST00390	421	EST00467
19	EST00023	122	EST00105	225	EST00178	330	EST00391	422	EST01447
21	EST00025	123	EST00106	226	EST00333	331	EST00392	423	EST00468
23	EST00026	125	EST00108	229	EST00180	332			
25		126	EST00109	231			EST00393	424	EST01448
27					EST00334	334	EST00395	425	EST00469
		127	EST00320	232	EST00182	335	EST00396	427	EST01449
28		129	EST00321	233	EST00183	337	EST00398	428	EST01451
29		130	EST00355	235	EST00185	340	EST00402	429	EST00471
. 30	EST00032	131	EST00322	236	EST00186	341	EST00403	431	EST00473
31	EST00033	133	EST00111	237	EST00187	342	EST00404	432	EST01452
32		134	EST00375	238	EST00188	344	EST00405		
33		135	EST00112	239	EST00189			434	EST00475
34						345	EST00406	435	EST00476
35		136	EST00113	240	EST00335	347	EST01829	436	EST00477
		138	EST00114	241	EST00191	348	EST01830	437	EST00478
36		139	EST00116	242	EST00192	349	EST01831	438	EST00479
39		140	EST00117	243	EST00193	350	EST00407	439	EST00480
- 40	EST00040	141	EST00118	244	EST00194	351	EST00408	440	EST01454
41	EST00041	142	EST00323	245	EST00347	352	EST00409	442	EST01456
42		143	EST00119	246	EST00196	353	EST00410		
46	EST00044	146	EST00122	250	EST00197			443	EST00482
• 47	EST00046	147				354	EST01433	444	EST00483
			EST00292	252	EST00198	355	EST00411	446	EST00485
49	EST00047	148	EST00236	254	EST00200	356	EST00412	447	EST00486
50	EST00048	149	EST00123	255	EST00201	357	EST00413	448	EST00487
51	EST00049	150	EST00124	256	EST00345	358	EST00414	449	EST00488
52	EST00052	151	EST00125	257	EST00337	359	EST00415	450	EST00489
53	EST00054	152	EST00126	259	EST00202	360	EST00416	451	
54	EST00055	153	EST00127	260	EST00357	361			EST00490
55	EST00056	154	EST00128	261			EST00417	452	EST00491
56					EST00338	363	EST00419	455	EST00494
	EST00057	155	EST00129	262	EST00339	364	EST00420	457	EST00495
57	EST00058	157	EST00131	265	EST00205	365	EST01434	458	EST00496
58	EST00059	158	EST00132	266	EST00206	366	EST00421	459	EST00497
. 59	EST00061	159	EST00325	272	EST00340	367	EST00422	460	EST01457
60	EST00062	160	EST00326	274	EST00268	369	EST00424	461	EST01836
63	EST00065	162	EST00133	275	EST00209	372	EST00427		
64	EST00066	163	EST00134	278	EST00342			462	EST00498
67	EST00351	165				373	EST01832	464	EST00499
			EST00136	279	EST00213	374	EST00428	465	EST00500
. 68	EST00068	167	EST00138	280	EST00343	375	EST00429	466	EST00501
69	EST00360	168	EST00140	283	EST00215	376	EST01436	467	EST00502
<u>71</u>	EST00070	169	EST00141	284	EST00216	377	EST00430	468	EST00503
73	EST00072	170	EST00295	286	EST00217	378	EST00431	470	EST00504
74	EST00073	171	EST00327	287	EST00218	379	EST00432	SEQ ID#	EST#
76	EST00075	172	EST00142	288	EST00219	380		<u> </u>	<u> </u>
80	EST00077	173	EST00143	289	EST00220		EST01439	177	ERTAREA/
81	EST00315	175	EST00144				EST00433	473	EST00506
83				290	EST00221	382	EST00434	474	EST00507
	EST00079	178	EST00294	291	EST00222	SEQ ID#	EST#	477	EST01463
84	EST00080	182	EST00329	292	EST00223			478	EST00510
85	EST00081	184	EST00149	293	EST00224	383	EST00435	479	EST00511
. 86	EST00082	185	EST00150	294	EST00225	384	EST01440	480	EST01464
87	EST00083	186	EST00151	SEQ ID#	EST#	386	EST00437	481	EST00512
89	EST00085	190	EST00153			388			
91	EST00086	191	EST00154	30F	ECT00224		EST00439	482	EST01465
92					EST00226	390	EST01442	483	EST00513
	EST00087	194	EST00157	297	EST00230	391	EST00441	484	EST00514
94	EST00353	SEQ ID#	EST#	298	EST00231	393	EST00443	487	EST00516
95	EST00088			302	EST00303	395	EST00445	488	EST00517
96	EST00089	195	EST00158	303	EST00348	397	E5:00446	489	EST00518
99	EST00316	196	EST00159	304	EST00307	398	EST00447	490	EST00519
SEQ ID#	EST#	197	EST00160	305	EST00308	399	EST00447		
			EST00161	306	EST00305	40C		491	ESTO0520
100	EST00090	199	EST00277				EST00449	492	EST00521
101				307		401	EST00450	495	EST00524
101	EST00091	203	EST00164	308	EST00314	403	EST00452	497	EST00526
									-

498		600	EST01492	- 697	EST00680	799	EST00752	894	EST01894
499	EST01468	601	EST01493	698	EST00681	800		895	
500	EST00527	603	EST01494	701	EST01522	801			EST01895
								896	EST01896
501		604	EST00607	702	EST00684	804	EST00757	897	EST01897
502	EST01469	605	EST00608	703	EST00685	805	EST00758	898	EST01898
507	EST00530	609	EST01496	704	EST00686	806			
508								899	EST01899
		610	EST00612	705	EST00687	807	EST00760	900	EST01900
509	EST01472	611	EST00613	706	EST00688	809	EST00762	901	EST01901
510	EST00532	612	EST00614	708	EST00689	810	,		
. 511	EST00533							. 902	EST01902
		615	EST00616	709	EST00690	811	EST00764	903	EST01903
512		616	EST01497	710	EST00691	813	EST00765	904	EST01904
513	EST00535	618	EST01498	711	EST00692	814	EST00766	905	
514		620							EST01905
			EST01499	712	EST00693	815	EST01855	906	EST01906
515	EST00537	622	EST01843	713	E\$T00694	816	EST01856	908	EST01908
519	EST00541	623	EST00621	714	EST00695	817		909	
520		624							EST01909
			EST01500	715	EST01523	818	EST01858	910	EST01910
521	EST01474	625	EST01844	716	EST01524	819	EST01859	911	EST01911
. 523	EST01838	627	EST00623	717	EST01525	820	EST01860	914	EST01914
525	EST00545	628	EST01503	718	EST00696				
527						822	EST01863	915	EST01915
		629	EST00624	719.	EST01526	825	EST01866	916	EST01917
528	EST00548	630	EST01505	720	EST00697	826	EST01867	917	EST01919
530	EST01477	. 631	EST00625	721	EST01527	827			
531		632					EST01558	918	EST01920
	EST00550		EST00626	722	EST01528	828	EST00767	920	EST01922
533	EST00552	633	EST00627	723	EST00698	830	EST01559	921	EST01923
534	EST01478	634	EST00628	725	EST00699	831	EST00769		
535	EST00553	636							EST01924
			EST01507	728	EST00702	832	EST00770	923	EST01925
536	EST01479	637	EST00630	730·	EST00704	837	EST01561	925	EST01927
537	EST00554	638	EST00631	731	EST00705	838	EST00774	926	EST01929
538	EST00555	640	EST01509	732					
					EST00706	839	EST01562	927	EST01930
539	EST00556	641	EST00633	733	EST00707	840	EST00775	928	EST01931
541	EST00558	643	EST00635	734	EST00708	841	EST00776	931	EST01934
542	EST01480	645	EST00637	735	EST00709				
543						842	EST01563	932	EST01935
	EST00559	646	EST00638	736	EST01532	843	EST01564	933	EST01936
544	EST00560	647	EST00639	737	EST00710	844	EST01565	934	EST01937
545	EST01481	648	EST00640	739	EST01534	846	EST00778	937	
547	EST00561	649	EST00641						EST01940
				740	EST01535	847	EST00779	939	EST01943
548	EST00562	651	EST00643	741	EST00712	848	EST01566	SEQ ID#	EST#
550	EST00564	652	EST01510	744	EST01537	849	EST01567		
553	EST00566	654	EST00644	746	EST00716			0/0	
						850	EST00780	940	EST01944
555	EST01483	655	EST00645	748	EST01850	851	E\$T00781	941	EST01945
556	EST00568	656	EST01513	749	EST00719	SEQ ID#	EST#	942	EST01947
558	EST01484	658	EST00647	750	EST01539	333 100			
560	EST01485						·	943	EST01948
		659	EST00648	751	EST01540	853	EST00783	· 944	EST01949
561	EST00571	661·	EST00650	754	EST00722	855	EST00785	945	EST01950
562	EST00572	. 662	EST00651	SEQ ID#	EST#	856	EST01568	946	EST01953
563	EST00573	663	EST00652	524 15#	<u> </u>				
				·		857	EST01868	947	EST01954
564	EST00574	664	EST00653	756	EST01541	858	EST01869	949	EST01958
565	EST00575	665	EST00654	758	EST00724	859	EST01870	950	EST01959
566	EST00576	SEQ ID#	EST#	761	EST01544	860			
567	EST00577	324 1511	LO117				EST00786	953	EST01962
				762	EST00727	861	EST01871	954	EST01963
568	EST00578	666	EST01514	763	EST00728	863	EST01873	956	EST01968
569	EST00579	667	EST00655	765	EST00730	864	EST00787	957	EST01969
SEQ ID#	EST#	668	.EST00656	766	EST00731	865	EST01569		
		7.7.2						958	EST01970
		669	EST00657	767	EST00732	866	EST01874	959	EST01972
571	EST00581	670	EST00658	768	EST00733	867	EST01875	960	EST01973
572	EST00582	671	EST00659	770	EST00735	868	EST01876	961	EST01974
574	EST00584	672	EST00660	771	EST01546				
575	EST00585		EST01515			869	EST00788	962	EST01975
		277					ECTIN7UN	963	EST01976
		673		772	EST00736	870	EST00789	. ,05	E31017/0
577	EST00587	673 674	EST01516	774	EST01548	870 871			
580		674	EST01516	.774	EST01548	871	EST00790	964	EST01977
580	EST00587 EST00590	674 675	EST01516 EST00661	774 775	EST01548 EST00737	871 872	EST00790 EST00791	964 966	EST01977 EST01979
580 581	EST00587 EST00590 EST00591	674 675 676	EST01516 EST00661 EST00662	.774 775 777	EST01548 EST00737 EST00739	871 872 873	EST00790 EST00791 EST00792	964 966 967	EST01977 EST01979 EST01980
580 581 583	EST00587 EST00590 EST00591 EST00593	674 675 676 677	EST01516 EST00661 EST00662 EST00663	774 775 777 779	EST01548 EST00737	871 872	EST00790 EST00791	964 966	EST01977 EST01979
580 581 583 584	EST00587 EST00590 EST00591	674 675 676	EST01516 EST00661 EST00662	774 775 777 779	EST01548 EST00737 EST00739 EST00741	871 872 873 874	EST00790 EST00791 EST00792 EST00793	964 966 967 970	EST01977 EST01979 EST01980 EST01983
580 581 583	EST00587 EST00590 EST00591 EST00593	674 675 676 677 678	EST01516 EST00661 EST00662 EST00663 EST01517	774 775 777 779 780	EST01548 EST00737 EST00739 EST00741 EST01549	871 872 873 874 875	EST00790 EST00791 EST00792 EST00793 EST00794	964 966 967 970 972	EST01977 EST01979 EST01980 EST01983 EST01986
580 581 583 584 585	EST00587 EST00590 EST00591 EST00593 EST00594 EST00595	674 675 676 677 678 679	EST01516 EST00661 EST00662 EST00663 EST01517 EST01518	774 775 777 779 780 781	EST01548 EST00737 EST00739 EST00741 EST01549 EST01550	871 872 873 874 875 876	EST00790 EST00791 EST00792 EST00793 EST00794 EST00795	964 966 967 970 972 974	EST01977 EST01979 EST01980 EST01983 EST01986 EST01988
580 581 583 584 585 586	EST00587 EST00590 EST00591 EST00593 EST00594 EST00595 EST00596	674 675 676 677 678 679 680	EST01516 EST00661 EST00662 EST00663 EST01517 EST01518 EST00664	774 775 777 779 780 781 783	EST01548 EST00737 EST00739 EST00741 EST01549 EST01550 EST01552	871 872 873 874 875 876 877	EST00790 EST00791 EST00792 EST00793 EST00794 EST00795 EST01877	964 966 967 970 972	EST01977 EST01979 EST01980 EST01983 EST01986
580 581 583 584 585 586 587	EST00587 EST00590 EST00591 EST00593 EST00594 EST00595 EST00596 EST01488	674 675 676 677 678 679	EST01516 EST00661 EST00662 EST00663 EST01517 EST01518	774 775 777 779 780 781	EST01548 EST00737 EST00739 EST00741 EST01549 EST01550	871 872 873 874 875 876	EST00790 EST00791 EST00792 EST00793 EST00794 EST00795 EST01877	964 966 967 970 972 974 975	EST01977 EST01979 EST01980 EST01983 EST01986 EST01988 EST01989
580 581 583 584 585 586	EST00587 EST00590 EST00591 EST00593 EST00594 EST00595 EST00596	674 675 676 677 678 679 680 682	EST01516 EST00661 EST00662 EST00663 EST01517 EST01518 EST00664 EST00666	774 775 777 779 780 781 783 785	EST01548 EST00737 EST00739 EST00741 EST01549 EST01550 EST01552 EST01553	871 872 873 874 875 876 877 878	EST00790 EST00791 EST00792 EST00793 EST00794 EST00795 EST01877 EST01878	964 966 967 970 972 974 975	EST01977 EST01979 EST01980 EST01983 EST01986 EST01988 EST01989 EST01990
580 581 583 584 585 586 587 588	EST00587 EST00590 EST00591 EST00593 EST00594 EST00595 EST00596 EST01488 EST00597	674 675 676 677 678 679 680 682 683	EST01516 EST00661 EST00662 EST00663 EST01517 EST01518 EST00664 EST00666 EST00667	774 775 777 779 780 781 783 785 786	EST01548 EST00737 EST00739 EST00741 EST01549 EST01550 EST01552 EST01553 EST00742	871 872 873 874 875 876 877 878	EST00790 EST00791 EST00792 EST00793 EST00794 EST00795 EST01877 EST01878 EST01879	964 966 967 970 972 974 975 976	EST01977 EST01979 EST01980 EST01983 EST01986 EST01988 EST01989 EST01990 EST01991
580 581 583 584 585 586 587 588 589	EST00587 EST00590 EST00591 EST00593 EST00594 EST00595 EST00596 EST01488 EST00597 EST00598	674 675 676 677 678 679 680 682 683	EST01516 EST00661 EST00662 EST00663 EST01517 EST01518 EST00664 EST00666 EST00667	774 775 777 779 780 781 783 785 786 787	EST01548 EST00737 EST00739 EST00741 EST01549 EST01550 EST01552 EST01553 EST00742 EST00743	871 872 873 874 875 876 877 878 879 880	EST00790 EST00791 EST00792 EST00793 EST00794 EST00795 EST01877 EST01878	964 966 967 970 972 974 975	EST01977 EST01979 EST01980 EST01983 EST01986 EST01988 EST01989 EST01990
580 581 583 584 585 586 587 588 589 590	EST00587 EST00590 EST00591 EST00593 EST00594 EST00595 EST00596 EST01488 EST00597 EST00598 EST00599	674 675 676 677 678 679 680 682 683 684 685	EST01516 EST00661 EST00662 EST00663 EST01517 EST01518 EST00664 EST00666 EST00667	774 775 777 779 780 781 783 785 786	EST01548 EST00737 EST00739 EST00741 EST01549 EST01550 EST01552 EST01553 EST00742	871 872 873 874 875 876 877 878	EST00790 EST00791 EST00792 EST00793 EST00794 EST00795 EST01877 EST01878 EST01879	964 966 967 970 972 974 975 976 977	EST01977 EST01979 EST01980 EST01983 EST01986 EST01988 EST01989 EST01990 EST01991 EST01992
580 581 583 584 585 586 587 588 589	EST00587 EST00590 EST00591 EST00593 EST00594 EST00595 EST00596 EST01488 EST00597 EST00598	674 675 676 677 678 679 680 682 683 684 685	EST01516 EST00661 EST00663 EST01517 EST01518 EST00664 EST00667 EST00668 EST00669	774 775 777 780 781 783 785 786 787 788	EST01548 EST00737 EST00739 EST00741 EST01550 EST01550 EST01552 EST00742 EST00744	871 872 873 874 875 876 877 878 879 880 882	EST00790 EST00791 EST00792 EST00793 EST00794 EST00795 EST01877 EST01877 EST01879 EST01880 EST01880	964 966 967 970 972 974 975 976 977 978 981	EST01977 EST01979 EST01980 EST01983 EST01986 EST01988 EST01989 EST01990 EST01991 EST01992 EST01995
580 581 583 584 585 586 587 588 589 590 591	EST00587 EST00590 EST00591 EST00593 EST00594 EST00596 EST00596 EST00597 EST00598 EST00599 EST00599 EST00489	674 675 676 677 678 679 680 682 683 684	EST01516 EST00661 EST00663 EST01517 EST01518 EST00664 EST00666 EST00667 EST00668 EST00669 EST00670	774 775 777 779 780 781 783 785 786 787 788	EST01548 EST00737 EST00739 EST00741 EST01549 EST01550 EST01552 EST00753 EST00743 EST00744 EST00745	871 872 873 874 875 876 877 878 879 880 882 883	EST00790 EST00791 EST00792 EST00793 EST00794 EST00795 EST01877 EST01878 EST01879 EST01880 EST01882 EST01883	964 966 967 970 972 974 975 976 977 978 981 982	EST01977 EST01979 EST01980 EST01983 EST01986 EST01989 EST01990 EST01991 EST01992 EST01995 EST01996
580 581 583 584 585 586 587 588 589 590 591 592	EST00587 EST00590 EST00591 EST00593 EST00594 EST00596 EST01488 EST00597 EST00598 EST00599 EST01489 EST00600	674 675 676 677 678 679 680 682 683 684 685 686	EST01516 EST00661 EST00662 EST00663 EST01517 EST01518 EST00664 EST00666 EST00667 EST00669 EST00670 EST00672	774 775 777 780 781 783 785 786 787 788 789 790	EST01548 EST00737 EST00739 EST00741 EST01550 EST01552 EST007553 EST00742 EST00743 EST00744 EST00745 EST00745	871 872 873 874 875 876 877 878 879 880 882 883 883	EST00790 EST00791 EST00792 EST00793 EST00794 EST00795 EST01877 EST01878 EST01879 EST01880 EST01882 EST01883 EST01885	964 966 967 970 972 974 975 976 977 981 981 982 983	EST01977 EST01979 EST01980 EST01983 EST01986 EST01988 EST01989 EST01990 EST01991 EST01992 EST01995
580 581 583 584 585 586 587 588 589 590 591 592 593	EST00587 EST00590 EST00591 EST00593 EST00594 EST00595 EST00597 EST00597 EST00598 EST00599 EST01489 EST00400 EST00601	674 675 676 677 678 679 680 682 683 684 685 686 688	EST01516 EST00661 EST00663 EST01517 EST01518 EST00664 EST00666 EST00667 EST00668 EST00669 EST00670	774 775 777 779 780 781 783 785 786 787 788	EST01548 EST00737 EST00739 EST00741 EST01549 EST01550 EST01552 EST00753 EST00743 EST00744 EST00745	871 872 873 874 875 876 877 878 879 880 882 883	EST00790 EST00791 EST00792 EST00793 EST00794 EST00795 EST01877 EST01878 EST01879 EST01880 EST01882 EST01883 EST01885	964 966 967 970 972 974 975 976 977 981 981 982 983	EST01977 EST01979 EST01980 EST01983 EST01986 EST01989 EST01990 EST01990 EST01991 EST01992 EST01995 EST01997
580 581 583 584 585 586 587 588 589 590 591 592	EST00587 EST00590 EST00591 EST00593 EST00594 EST00596 EST01488 EST00597 EST00598 EST00599 EST01489 EST00600	674 675 676 677 678 679 680 682 683 684 685 686	EST01516 EST00661 EST00663 EST01517 EST01518 EST00664 EST00666 EST00667 EST00668 EST00670 EST00672 EST00672	774 775 777 779 780 781 783 785 786 787 788 789 790	EST01548 EST00737 EST00739 EST00741 EST01549 EST01552 EST01552 EST00742 EST00743 EST00744 EST00745 EST00745	871 872 873 874 875 876 877 878 880 882 883 885 885	EST00790 EST00791 EST00792 EST00794 EST00795 EST01877 EST01878 EST01889 EST01882 EST01883 EST01885 EST01887	964 966 967 970 972 974 975 976 977 981 981 982 983 984	EST01977 EST01979 EST01980 EST01980 EST01986 EST01988 EST01989 EST01990 EST01991 EST01991 EST01995 EST01996 EST01997 EST01998
580 581 583 584 585 586 587 589 590 591 592 593	EST00587 EST00590 EST00591 EST00593 EST00594 EST00595 EST00597 EST00597 EST00598 EST00599 EST01489 EST00600 EST00601 EST01840	674 675 676 677 678 679 680 682 683 684 685 686 688	EST01516 EST00661 EST00663 EST01517 EST01518 EST00664 EST00667 EST00669 EST00669 EST00672 EST00672 EST00674 EST00676	774 775 777 779 780 781 783 785 786 787 788 789 790 792	EST01548 EST00737 EST00739 EST00741 EST01550 EST01550 EST01553 EST00742 EST00743 EST00744 EST00745 EST00745 EST00747 EST00748	871 872 873 874 875 876 877 878 879 880 882 883 885 887 389	EST00790 EST00791 EST00792 EST00793 EST00794 EST01877 EST01878 EST01880 EST01880 EST01882 EST01883 EST01887 CST01887 CST01887	964 966 967 970 972 974 975 976 977 978 981 982 983 983	EST01977 EST01979 EST01980 EST01983 EST01986 EST01988 EST01990 EST01991 EST01991 EST01995 EST01995 EST01997 EST01998 EST01998
580 581 583 584 585 586 587 588 589 590 591 592 593 593	EST00587 EST00590 EST00591 EST00593 EST00594 EST00595 EST00596 EST00597 EST00598 EST00599 EST01489 EST00600 EST00601 EST01840 EST00607	674 675 676 677 678 679 680 682 683 684 685 686 688 690 692	EST01516 EST00661 EST00663 EST01517 EST01518 EST00664 EST00667 EST00667 EST00669 EST00670 EST00670 EST00674 EST00674	774 775 777 779 780 781 783 785 786 787 788 789 790 792 793	EST01548 EST00737 EST00739 EST00741 EST01550 EST01552 EST01553 EST00742 EST00743 EST00744 EST00745 EST00745 EST00747 EST00748 EST00748	871 872 873 874 875 876 877 878 879 880 882 883 885 887 389	EST00790 EST00791 EST00792 EST00793 EST00794 EST00795 EST01877 EST01877 EST01878 EST01880 EST01882 EST01883 EST01885 EST01887 CST01889 EST01889	964 966 967 970 972 974 975 976 977 981 982 983 984 987	EST01977 EST01979 EST01983 EST01986 EST01986 EST01989 EST01990 EST01991 EST01992 EST01995 EST01996 EST01997 EST01998 EST02001 EST02003
580 581 583 584 585 586 587 588 589 590 591 592 593 595 597	EST00587 EST00590 EST00593 EST00594 EST00596 EST00596 EST01488 EST00597 EST00598 EST00599 EST01489 EST00600 EST00601 EST01840 EST00604	674 675 676 677 678 680 682 683 684 685 686 688 690 692 693 694	EST01516 EST00661 EST00663 EST01517 EST01518 EST00664 EST00667 EST00669 EST00670 EST00670 EST00674 EST00674 EST00676 EST00677 EST00677	774 775 777 779 780 781 783 785 786 787 788 789 790 /92 793	EST01548 EST00737 EST00737 EST00741 EST01550 EST01552 EST01552 EST00742 EST00743 EST00744 EST00745 CCT01554 EST00748 EST00748 EST00748 EST00750	871 872 873 874 875 876 877 878 879 880 882 883 885 887 389	EST00790 EST00791 EST00792 EST00793 EST00794 EST01877 EST01878 EST01880 EST01880 EST01882 EST01883 EST01887 CST01887 CST01887	964 966 967 970 972 974 975 976 977 978 981 982 983 983	EST01977 EST01979 EST01980 EST01983 EST01986 EST01988 EST01990 EST01991 EST01991 EST01995 EST01995 EST01997 EST01998 EST01998
580 581 583 584 585 586 587 588 589 590 591 592 593 593	EST00587 EST00590 EST00591 EST00593 EST00594 EST00595 EST00596 EST00597 EST00598 EST00599 EST01489 EST00600 EST00601 EST01840 EST00607	674 675 676 677 678 679 680 682 683 684 685 686 688 690 692	EST01516 EST00661 EST00663 EST01517 EST01518 EST00664 EST00667 EST00667 EST00669 EST00670 EST00670 EST00674 EST00674	774 775 777 779 780 781 783 785 786 787 788 789 790 /92 793	EST01548 EST00737 EST00737 EST00741 EST01550 EST01552 EST01552 EST00742 EST00743 EST00744 EST00745 CCT01554 EST00748 EST00748 EST00748 EST00750	871 872 873 874 875 876 877 878 879 880 882 883 885 887 389 890	EST00790 EST00791 EST00792 EST00793 EST00794 EST00795 EST01877 EST01878 EST01880 EST01880 EST01883 EST01885 EST01887 CST01887 EST01889 EST01890 EST01892	964 966 967 970 972 974 975 976 977 978 981 982 983 984 987	EST01977 EST01979 EST01980 EST01983 EST01986 EST01989 EST01990 EST01991 EST01992 EST01995 EST01996 EST01997 EST01997 EST01997 EST01998 EST01997 EST01998
580 581 583 584 585 586 587 588 589 590 591 592 593 595 597	EST00587 EST00590 EST00593 EST00594 EST00596 EST00596 EST01488 EST00597 EST00598 EST00599 EST01489 EST00600 EST00601 EST01840 EST00604	674 675 676 677 678 680 682 683 684 685 686 688 690 692 693 694	EST01516 EST00661 EST00663 EST01517 EST01518 EST00664 EST00667 EST00669 EST00670 EST00670 EST00674 EST00674 EST00676 EST00677 EST00677	774 775 777 779 780 781 783 785 786 787 788 789 790 792 793	EST01548 EST00737 EST00739 EST00741 EST01550 EST01552 EST01553 EST00742 EST00743 EST00744 EST00745 EST00745 EST00747 EST00748 EST00748	871 872 873 874 875 876 877 878 879 880 882 883 885 887 389	EST00790 EST00791 EST00792 EST00793 EST00794 EST00795 EST01877 EST01877 EST01878 EST01880 EST01882 EST01883 EST01885 EST01887 CST01889 EST01889	964 966 967 970 972 974 975 976 977 981 982 983 984 987	EST01977 EST01979 EST01983 EST01986 EST01986 EST01989 EST01990 EST01991 EST01992 EST01995 EST01996 EST01997 EST01998 EST02001 EST02003

992	EST02006	1086	EST02105	1184	EST02209	1274	EST02303	1363	ECTO370/
994	EST02008	1087	EST02106	1185	EST02210	1275	EST02304		EST02394
997	EST02011	1088						1364	EST02395
999			EST02107	1186	EST02211	1276	EST02305	1366	EST02397
	EST02013	1091	EST02110	1187	EST02212	1278	EST02307	1367	EST02398
1001	EST02015	1093	EST02112	1188	EST02213	1279	EST02308	1368	EST02399
1002	EST02016	1095	EST02114	1189	EST02214	1280	EST02309	1370	EST02401
1003	EST02017	1097	EST02116	1190	EST02215	1281	EST02310	1372	EST02403
1005	EST02019	1098	EST02117	1191	EST02216	1282	EST02311	1373	EST02404
1006	EST02020	1100	EST02119	1192	EST02217	1283			
1008	EST02022	1101					EST02312	1375	EST02406
1009			EST02120	1193	EST02218	1284	EST02313	1376	EST02407
	EST02023	1102	EST02121	1194	EST02219	1286	EST02316	1377	EST02408
1010	EST02024	1104	EST02123	1195	EST02220	1288	EST02318	1378	EST02409
1011	EST02025	1106	EST02125	1197	EST02222	1289	EST02319		EST02410
1012	EST02026	1107	EST02126	1198	EST02223	1290	EST02320		EST02411
1013	EST02027	1108	EST02127		EST02224	1291	EST02321	1381	
1014	EST02028	1109	EST02128		EST02226	1292			EST02413
1015	EST02029	1110	EST02129	1201			EST02322	1382	EST02414
					EST02228	1293	EST02323		
1016	EST02030	1111	EST02131		EST02229	1294	EST02324		
	EST02031	1112	EST02132	1203	EST02230	1295	EST02325		
1019	EST02033	1114	EST02134	1204	EST02232	1296	EST02326		
1022	EST02036	1117	EST02137	1206	EST02234	SEQ ID#	EST#		
1023	EST02037	1119	EST02139	1207	EST02235		<u> </u>		
1024	EST02038	1120	EST02140	1208	EST02236	1298	EST02328		
1025	EST02040	1121	EST02141	1209		1299	EST02329		
1026	EST02040	1122							
			EST02142	SEQ ID#	EST#	1300	EST02330		
1027	EST02042	1123	EST02143	45		1302	EST02332		
1028	EST02043	1124	EST02144	1211	EST02239	1303	EST02333		
· 1029	EST02044	1125	EST02145	1212	EST02240	1304	EST02334		•
1030	EST02045	SEQ ID#	EST#	1213	EST02241	1305	EST02335		
1032	EST02048			1214	EST02242	1306	EST02336		
1033	EST02049	1127	EST02147	1215	EST02244	1307	EST02337		. •
	EST02052	1128	EST02148	1216	EST02245	1307	EST02339		
					E3102243				
SEQ ID#	EST#	1130	EST02150	1217		1310	EST02340		
		1131	EST02151	1218	EST02247	1311	EST02341		
1037	EST02053	1132	EST02152	1219	EST02248	1313	EST02343		
1038	EST02054	1135	EST02155	1220	EST02249	1314	EST02344		
1040	EST02056	1136	EST02156	1221	EST02250	1315	EST02345		
. 1042	EST02058	1137	EST02157		EST02252	1316	EST02346	•	
1044	EST02060	1138	EST02159	1225	EST02254	1317	EST02347		
1045	EST02061	1140	EST02162	1226	EST02255	1318	EST02348	•	
	EST02062	1142	EST02164	1227	EST02256				
1048						1319	EST02349		
	EST02064	1143	EST02165	1232	EST02261	1320	EST02350		
1049	EST02065	1144	EST02166	1234	EST02263	1321	EST02351		
1050	EST02066	1145	EST02167	1235	EST02264	1322	EST02352		
. 1051	EST02067	1148	EST02170	1236	EST02265	1323	EST02353		
1052	EST02068	1149	EST02171	1237	EST02266	1325	EST02355		
1053	EST02069	1150	EST02172		EST02267	1326	EST02356		
1054	EST02070	1152	EST02174		EST02268	1327	EST02357		
1055	EST02071	1153	EST02175	1240	EST02269	1328	EST02358		
1056	EST02072	1154	EST02176		EST02270	1329	EST02359		
1057	EST02073			1241	ES102270	1327			
		1155	EST02177		EST02271	1330	EST02360		
1058	EST02074	1156	EST02178	1244	EST02273	1333	EST02363		
1059	EST02075	1157	EST02180		EST02275	1334	EST02364		
1060	EST02076 -	1158	EST02181	1247	EST02276	1335	EST02365		
1061	EST02078		EST02182		EST02277	1336	EST02366		
1062	EST02079	1160	EST02183	1249	EST02278	1337	EST02367		
1063	EST02081	1161	EST02184		EST02279	1338	EST02368		
1064	EST02082	1162	EST02185		EST02280	1339	EST02369		
1065	EST02083	1164	EST02188		EST02281	1342	EST02372		
1066	EST02084		EST02189	1257	EST02282	1343			
1067	EST02085			123	FOTOSSOZ		EST02373		
			EST02190	1234	EST02283	1345	EST02375		
1068	EST02086		EST02191		EST02284	1346	EST02376		
	EST02088		EST02193		EST02285	1347	EST02377		
1071	EST02089		EST02194		EST02286	1349	EST02379		
	EST02090	1170	EST02195	1258	EST02287	1350	EST02380		
	EST02091	1171	EST02196	1259	EST02288	1351	EST02381		
	EST02092	1172	EST02197	1260	EST02289	1352	EST02382		
1075	EST02093	1173	EST02198		EST02290	1353	EST02383		•
	EST02094	1174	EST02199		EST02290	1354	EST02384		
	EST02096	1175	EST02200	1203	EST02292	1355	EST02385		
	EST02097	1176	5ST02201	1208	EST02297	1357	EST02387		
1079	EST02098	1177	EST02202	1269	EST02298	1358	EST02388		
1080	EST02099	1178	EST02203	1270	EST02299	1359	EST02390		
1082	EST02101	1179	EST02204	1271	EST02300	1360	ESTO2391		\$4.TT
1084	EST02103	1180	EST02205	1272	EST02301	1361	EST02392		
1085	EST02104	1182	EST02207	1273		1362	EST02393		
					_				

SEQ ID#	EST#	1485	EST02522	1592	EST02637	1689	EST00845	1799	EST00934
		1486	EST02523	1593		1690		1800	
1384	EST02416	1487		1594	EST02639	1691		1801	EST00936
1387	EST02419	1488		1596	EST02641	1696		1802	
1389	EST02422	1489		1597		1697	EST00852	1803	
1390	EST02423	1490		1598	EST02643	1702		1806	
1391	EST02424	1491		1600	EST02645	1703		1808	
1392	EST02425	1494		1601		1705		1810	
1393	EST02426	1497		1603		1707	EST01581	1812	EST02693
1394	EST02427	1498		1604	EST02649	1709		1813	EST00946
1396	EST02430.	1501		1605	EST02650	1711	EST00861	1814	EST00947
1398	EST02432	1504		1606		1712		1815	EST01615
1400	EST02434	1506		1607	EST02652	1713		1816	EST00948
1402	EST02436	1507		1609		1714		1817	EST00949
1403	EST02437	1508		1611	EST02656	1717	EST00866	1818	EST01616
1404	EST02438	1509		1612	EST02657	1719	EST00868	1821	EST00952
1406	EST02440.	1510		1613	EST02658	1720	EST00869	1822	EST00953
1407	EST02441	. 1512		1614	EST02659	1721	EST00870	1823	EST00954
1410	EST02444	. 1513	EST02552	1615	EST02660	1722	EST00871	1824	EST01617
1411	EST02445	1514	EST02553	1617	EST02662	1723	EST00872	1825	EST00955
1414	EST02448	1515	EST02554	1618	EST02663	1724	EST00873	1827	EST01618
1415	EST02449	1517		1619	EST02665	1725	EST00874	1828	EST00957
1416	EST02450	1518	EST02559	1620	EST02666	1727	EST00875	1831	EST01619
1419	EST02454	1519	EST02560	1622	EST02668	1728	EST00876	1832	EST00960
1420	EST02456	1520	EST02561	1623	EST02669	1732	EST01590	1833	EST00961
1421	EST02457	1521	EST02562	1625	EST02672	1733	EST01591	1835	EST00962
1422	EST02458	1522	EST02563	1626	EST02673	1734	EST00880	1836	EST01622
1423	EST02459	1523	EST02564	1628	EST02675	1735	EST00881	1837	EST00963
1424	EST02460	1524	EST02565	1632	EST02679	1736	EST01592	1838	EST00964
	EST02461	1525	EST02566	1633	EST02680	1737	EST00882	1839	EST00965
1426	EST02462	1526	EST02567	1634	EST02681	1740	EST02687	1840	EST00966
1428	EST02464	1529	EST02570	1635	EST02682	1741	EST00886	1841	EST00967
	EST02465	1530	EST02571	1636	EST02684	1744	EST00889	1842	EST00968
1431	EST02467	1532	EST02573	1638	EST00798	1745	EST00890	1843	EST00969
1432	EST02468	1533	EST02574	1640	EST00800	1747	EST00892	1844	EST00970
1433	EST02469	1534	EST02575	1641	EST00801	1748	EST00893	1846	EST00972
1434	EST02470	1535	EST02576	1642	EST00802	1749	EST01593	1850	EST01624
1435	EST02471	1538	EST02579	1643	EST00803	1750	EST00894	1851	EST00976
1436	EST02472	1539	EST02580	1645	EST00804	1752	EST00896	1854	EST00978
1437	EST02473	1540	EST02581	1646	EST00805	1753	EST00897	. 1855	EST00979
1438 1440	EST02474	1541	EST02582	1647	EST00806	1754	EST00898	1857	EST00980
1442	EST02476	1542	EST02583	1648	EST00807	1755	EST00899	1858	EST00981
1443	EST02478	1545	EST02587	1650	EST00809	1757	E\$T01594	1859	EST00982
1444	EST02479 EST02480	1546	EST02588	1652	EST00811	1758	EST00901	1861	EST00984
1445	EST02480	1547	EST02589	1653	EST00812	1759	EST00902	1862	EST00985
	EST02481	1549	EST02591	1655	EST00813	1761.		1863	EST00986
1447	EST02483	1550 1552	EST02592	1656	EST00814	1762	EST00904	1864	EST00987
	EST02486	1553	EST02594 EST02595	1657	EST00815	1763	EST00905	1865	EST00988
1452	EST02488	1554	EST02597	1658	EST00816	1765	EST01600	1867	EST00990
1453	EST02489	1555	EST02598	1659	EST00817	1766	EST00906	1868	EST00991
1454	EST02490	1556	EST02599	1660	EST00818	1772	EST02691	1870	EST00993
1455	EST02491	1557	EST02600	1661	EST00819	1773	EST00911		EST00995
1456	EST02492	1559	EST02602	1663	EST00820 EST00821	1774	EST00912	1873	EST01630
1458	EST02495	1560	EST02603			1775	EST02692	1874	EST00996
1459	EST02496	1564	EST02607	1664 1665	EST00822 EST00823		EST01603	1875	EST01631
1460	EST02497	1565	EST02608		EST00823	1778	EST00914	1876	EST00997
1461	EST02498	1568	EST02611	1668		1779	EST00915	SEQ ID#	EST#
1462	EST02499	1569	EST02612	1669	EST00826 EST00827	1781	EST00917		
1464	EST02501	1570	EST02613	1670	EST00828	1782	EST00918	1878	EST00999
1466	EST02503	1571	EST02614			1783	EST00919	1879	EST01633
1467	EST02504	1573	EST02616	1671 1672	EST00829	SEQ ID#	EST#	1881	EST01000
1469	EST02506	1574	EST02617	1673	EST00830	170/	F0*00000	1882	EST01638
1470	EST02507	1576	EST02619	1674	EST00831 EST00832	1784	EST00920	1883	EST01001
1471	EST02508	1577	EST02620			1785	EST00921	1884	EST01002
1472	EST02509	1578	EST02621	SEQ ID#	EST#	1786	EST00922	1886	EST01003
1474	EST02511	1579		1675	ECTODOTE	1787	EST00923	1887	EST01004
1475	EST02512	1580	EST02623		EST00833	1788	EST00924	1889	EST01006
1476	EST02512	SEQ ID#	EST#	1676	EST00834	1789	EST00925	1891	EST01008
1477	EST02514	1D#		1678 1679	EST00836	1790	EST00926	1893	EST01642
1481	EST02518	1582	EST02626	1680	ESTOOB37	1791	EST00927	1895	EST01010
1482	EST02519	1583	EST02628	1684	EST00838	1792	EST00928	1898	EST01013
SEQ ID#	EST#	1584	EST02629	1685	EST00841	1794	EST01607	1899	EST01014
		1585	EST02630		EST00842 EST01574	1795 1796	EST00930	1901	EST01016
1483	EST02520		EST02632	1687	EST00843		EST00931		EST01017
1484	EST02521	1590	EST02635	1688	EST00844	1797	EST00932	1905	EST01020
		.570		1000	23100044	1798	EST00933	1906	EST01021

190		2016	5 EST01110	2118	B EST01194	2223	S EST01742	2332	5070170/
190		2018	3 EST01111	2119		222		2333	EST01794 EST01357
190		2019	P EST01112	212			EST01280	2335	EST01357
191		2020		2123	5 EST01713	2229	EST01281	2336	EST01360
191		2021		2124	EST01198	2231		2337	EST01361
1913		2022		2125		2237		2340	EST01802
191		2023		2126	S EST01200	2238		2341	EST01364
1916		2025		2127		2239		2343	EST01366
1917		2026		2129	EST01203	2240		2344	EST01367
1918		2027		2130	EST01204	2241	EST01747	2349	EST01372
1919		2028		2132	EST01206	2242	EST01292	2350	EST02708
1920		2029		2133	EST01207	2243	EST01293	2352	EST01374
1921		2030		2135	EST01209	2244		2356	EST01377
1922		2033		2137	' EST01211	2246		2357	EST01378
1923		2034		2139		2247		2360	EST01381
1924		2035		2140		2249		2361	EST01382
1925		2036		2142		2251	EST01300	2363	EST01384
1926		2037		2143		2252	EST01750	2364	EST01385
1927		2038		2147		2253	EST01301	2365	EST01386
1929		2039		2148		2256	EST02718	2366	EST01387
1932 1934		2040		2151		2257	EST01303	2369	EST01811
1934		2042		2152		2258	EST01754	2370	EST01390
		2045		2154		2260	EST01305	2371	EST01391
1936 1937		2047		2156		2261	EST01755	2372	EST01392
1938		2048		2157		2262	EST01306	2375	EST01815
1941		2049		2158		2264	EST01308	2376	EST01395
1941		2050		2159		2265	EST01309	2377	EST01396
1942		2052		2160		2268	EST01311	2379	EST01398
1945		2053 2054	EST01140	2162	EST01232	2269	EST01312	2380	EST01399
1946		2055	EST01141		EST01233	2270	EST01313	2381	EST01400
1947			EST01690 EST01143	2164		2271	EST01314	2382	EST01401
1948		2057		2165	EST01720	2272	EST01762	2383	EST01402
1950		2062	ES101147 EST02701	2166		2273	EST01315	2384	EST01403
1951	EST01056	2063	EST01148	2169	EST01237	2275	EST01316	2385	EST01816
1952	EST01057	2065	EST01691		EST01722	2276	EST01317	2386	EST01404
1955	EST01662	2066	EST01692	2171		2277	EST01318	2387	EST01405
1957	EST01059	2067	EST01693	2172	EST01240 EST01241	2278	EST01319		•
1958	EST01060	2069	EST01150	2175	EST01243	2279 2280	EST01320		
1959	EST01061	2070	EST01151	2177		2284	EST01763		
1963	EST01063	2072	EST01152	2178	EST01726		ESTO1323		
1964	EST01064	2074	EST01698	2179	EST01246	SEQ_ID#	EST#		
1966	EST01065	2075	EST01153	2180	EST01247	2285	EST01768		
1968	EST01067	2076	EST02702	2181	EST01248	2287	EST01770	•	
1969	EST01068	2077	EST01154	SEQ ID#	EST#	2288	EST01324		
1970	EST01666	2078	EST01155			2290	EST01772		
1971	EST01069	2079	EST01156	2182	EST01249	2291	EST01773		•
1972	EST01070	2080	EST01157	2183	EST01250	2292	EST01326		
. 1975	EST01073	SEQ ID#	EST#	2185	EST01252	2293	EST01327		
1976	EST01074			2186	EST01253	2294	EST01328		
· 1978	EST01076	2081	EST01158	2187	EST01727	2295	EST01329		
1979	EST01077	2082	EST01159 ·	2188	EST01254	2296	EST01330		
SEQ ID#	EST#	2083	EST01160	2190	EST01728	2298	EST01331		
1020	EST01078	2084	EST01161	2191	EST01256	2299	EST01332		
1981	EST01079	2085	EST01162	2193	EST01258	2301	EST01334		
1983	EST01079	2086 2087	EST01163	2194	EST01729	2304	EST01780		
1984	EST01082	2088	EST01164 EST01166	2195	EST01259	2305	EST01336		
1985	EST01083	2091	EST01168	2197	EST01261	2306	EST01337		
1986	EST01084	2093	EST01170	2198 2199	EST01730	2310	EST01341		
1988	EST01085	2095	EST01701		EST01262	2311	EST01342		
1989	EST01086	2096	EST01701	2200 2201	EST01731	2312	EST01343		
1995	EST01092	2097	EST01172	2202	EST01263 EST01732	2313 2315	EST01344		•
1996	EST01093	2098	EST01174	2205	EST01735	2316	EST01346 EST01782		
1998	EST01095	2099	EST01175	2206	EST01736	2317	EST01762		
1999	EST01096	2103	EST01179	2208	EST01267		EST01347		
2002	EST01099	2104	EST01180	2209	EST02717		EST01349		
2003	EST01675	2107	EST01183	2210	EST01268	2321	EST01350		
2005	EST01100	2108	EST01184	2211	EST01269	2322	EST01351		
2006	EST01101	2109	EST01185	2213	EST01271	2323	EST01789		
2007	EST01102		EST01186	2215	EST01273	2325	2ST01353		
2009	EST01677	2111	EST01187	2218	EST01274		EST01354		
2010	EST01104	2112	EST01188	2219	EST01275		EST01355		
	EST01105	2113	EST01189	2220	20701740	2329	EST01792		gar ay
2014	EST01108	2114	EST01190	2221	EST01741	2330	EST01793		•
2015	EST01109	2115	EST01191	2222	EST01276	2331	EST01356		

SEQ ID#	EST#
2389	EST01407
2391	EST01415
2392	EST01416
2395	EST01419
2397	EST01421
2401	EST01424
2403	EST01425
2404	EST01426
2406	EST02713
2409	EST00273

10

15

20

, 123775 - 123775

EXAMPLE 10

Functional Groupings of ESTs and Corresponding Genes

By matching new human ESTs to known sequences from other species, the apparent function of the gene corresponding to the EST can be ascertained. The data generated in Example 3 and 4 have been used to categorize 127 of the ESTs of the present invention, and their corresponding genes, into predicted functional groups. (These 127 are ESTs with database matches to sequences from other species for which a function was known.) Two different grouping schemes have been used.

The first scheme separates the sequences into three broad categories: metabolic; regulatory; and structural. These groupings are set out in Table 10.

The second grouping scheme separates the sequences into 13 specific categories: cell surface proteins; developmental control; energy metabolism; kinases and phosphatases; oncogenes; other metabolism-related polypeptides; peptidases and peptidase inhibitors; receptors; structural and cytoskeletal; signal transduction; transporters; transcription, translation, and subcellular localization; and transcription factors. These groupings are set out in Table 11.

Table 10: Thre -Class Functional Groupings of ESTs

SEQ ID	EST#	Group	Putative Identification
1834	EST01620	м	AMD doomings busin
97	EST00289	M M	AMP deaminase, brain Aconitase
	EST00233	M	Alcohol dehydrogenase
	EST01700	M	Anion exchanger homolog AE3
	EST01443	M	CDPdiacylglycerol-serine O-phosphatidyltransfera
	EST01663	M	Ca2+-transporting ATPase 2
	EST02055	M	Calcium channel
	EST01257		Diacylglycerol kinase, lymphyocyte
	EST02477	M	Diamine acetyltransferase
2289	EST01325	M	Fatty acid synthase
	EST01323	M	Fo ATPase beta subunit, mitochondrial
	EST00825	M	Gamma-aminobutyric acid transporter
	EST02446	M	Glutamate-aspartate carrier protein
	EST02034	М	Glutaminase
	EST01791	M	Inositol-1,4,5-trisphosphate 3-kinase
	EST01731		Lon protease
	EST02463	M	Long-chain-fatty-acid-CoA ligase
,	EST01744	M	NAD(P) + transhydrogenase (B-specific)
	EST02609	M	Neutrophil oxidase factor
	EST01573	M	
•	EST01751	M	Nucleoside diphosphate kinase
	EST00287	M	Phosphatidylinositol-4,5-bisphosphate phosphodie
	EST01775	M	Processing enhancing protein
	EST01775	M	Prohormone cleavage enzyme
	EST01572	M	Prolyl endopeptidase
	EST00374		Protochlorophyllide reductase
	EST01583	M	RNA polymerase II 6th subunit (RPO26) Ribosomal protein L18a
	EST01583		
	EST01627	M	Ribosomal protein Lla
	EST00300	M	Ribosomal protein L3
	EST00300	M	Ribosomal protein L30
	EST01826	M	Ribosomal protein \$10 Ribosomal protein \$10
	EST01828	M	
	EST01697		Ribosomal protein YL10 Succinate dehydrogenase flavoprotein
	EST01715	M	Succinate dehydrogenase flavoprotein
	EST01601	M	Thiosulfate sulfurtransferase (rhodanese)
	EST01711	M	Valine-tRNA ligase
	EST01588	M	XPR2 alkaline extracellular protease
	EST01913	M	Clathrin coat assembly protein AP50 homolog
	EST02051	M	J1 protein
	EST01982	R	ADP-ribosylation factor 1
	EST02146	R	Calbindin D28
1910	EST01645	R	Calmodulin
485	EST01466	R	Calmodulin-dependent protein kinase, type II, be
2302	EST01779	R	Discs-large tumor suppressor
188	EST00256	R	Enhancer of split
1229	EST02258	R	KUP protein
	EST02007		Kinase 5 protein
2282	EST01764	R	Lamin B receptor
SEQ ID	EST#	Group	
529 15	201H	oroup.	racactve luciteticación
161	EST00247	R	MARCKS (myristoylated alanine-rich protein kinas
769	EST00734	R R	MARCKS homolog
	EST02418	R	MARCKS homolog
	EST00259	R	Notch/Xotch
	EST01253	R	Notch/Xotch
	EST02429	R	Nuclear factor 1-like protein (NF1)
	EST01806	R	Prohibitin
4	EST02087	R	Protein kinase C, zcta
1933	EST01650	R	Protein phosphatase 2A beta subunit
~		•• (buophingane su seed ammitte

```
202 EST00298 R
                         Protein-tyrosine phosphatase LRP
  1478
       EST02515
                 R
                         Rab5
                         Seven in absentia
  1408
       EST02442
                 R
   300 EST00232
                  R
                         Transforming protein (dbl)
  1147 EST02169
                 R
                         Tyrosine kinase
  1348 EST02378
                 R
                         cAMP-dependent protein kinase inhibitor
  1931 EST01041
                         cAMP-regulated phosphoprotein
  1413 EST02447
                 R
                         cAMP-specific phosphodiesterase
                         ras p21-like small GTP-binding protein (smg GDS)
    37
       EST00038
                 R
   102
       EST00248
                 R
                         rho H12/ ARH12
   299 EST00249
                 R
                         smg p25A GDP dissociation inhibitor
   189 EST00282
                         trkB
  1332 EST02362
                         GA binding protein, beta subunit
  1277
       EST02306
                 R
                         Bib protein
    43
       EST00371
                 R
                         Maternal G10 protein
  1704
       EST01580
                         Myeloid differentiation primary response gene My
                 R
                         Otd homeotic protein
   346 EST01828
   187
       EST00152
                         Wilm's tumor-related protein
   249
       EST00275
                 R
                         Zinc Finger Proteins
   413
       EST01446
                 R
                         Zinc Finger Proteins
   469 EST01460
                         Zinc Finger Proteins
                 R
                         Zinc Finger Proteins
   833 EST01560
                 R
 1230 EST02259
                 R
                         Zinc finger proteins
 1496 EST02534
                 R
                         Zinc finger proteins
  2324
       EST01352
                         Zinc Finger Proteins
                 R
                         60K filarial antigen
   208
       EST00250
                 S
                         60K filarial antigen
 2320
       EST01784
                 S
   251
       EST00370
                 S
                         Actin, other
 2146 EST01218
                 S
                       Actin, other
                 S
   248
       EST00271
                        Actinin, alpha
       EST01891
   891
                        Actinin, alpha
                 S
 1500 EST02538 S
                        Actinin, alpha
  132 EST00110
                 S
                         Agrin
 1852
       EST01625
                 s
                        Agrin
 1965
       EST01664
                 S
                        Amyloid A4
                        Amyloid A4
 2068
       EST01694
                 S
 2408
       EST00244
                        Amyloid A4
 1880
       EST01634
                 S
                        Axonal glycoprotein TAG-1
 2004
       EST01676
                 S
                         Cofilin
   650
       EST00642
                 S
                        Dilute (myosin heavy chain)
 2217
       EST01738
                 S
                        Gelation factor ABP-280
 1885 EST01639
                        Histocompatibility antigen modifier 1
       EST00257
                        Kinesin
   77
                 S
SEQ ID
       EST#
                 Group
                        Putative Identification
   78 · EST00258
                         Kinesin
 2245 EST01748
                         Kinesin
  313 EST00276
                 S
                        Lysosomal membrane glycoprotein 1 (LAMP-1)
   223 EST00368
                 S
                        Microtubule-associated protein 1B
  824
       EST01865
                 S
                        Microtubule-associated protein 1B
 2032 EST01683
                        Microtubule-associated protein 1B
                 S
 2017 EST01678
                        Milk fat globule membrane protein
 1567 EST02610
                        Neural cell adhesion molecule L1
                 S
  506
       EST01471
                 S
                        Neuraxin
 2368
       EST01389
                 S
                         Radial spoke protein 3
                         Spectrin, beta
  951
       EST01960
                 S
 2089
       EST01699
                         Sperm membrane protein
                 S
   653
       EST01512
                         Tubulin, alpha
       EST00270
                 S
  311
                         Tubulin, beta
   594
       EST01490
                         Tubulin, beta
  757
       EST01542
                 S
                         Tubulin, beta
       EST02274 S
 1245
                         Tubulin, beta
 1589
       EST02634 S
                         Tubulin, beta
 1468 EST02505 S
                        Matrin 3
```

-70-

1371 EST02402 S Talin 1701 EST00853 S Unc-104

Group Key: M: Metabolic, R: Regulatory, S: Structural

Table 11: Thirteen-Class Functional Groupings of ESTs

SEQ ID	EST#	Group	Putative Identification
208	EST00250	CS	60K filarial antigen
2320	EST01784	CS	60K filarial antigen
1965	EST01664	CS	Amyloid A4
2068	EST01694	CS	Amyloid A4
2408	EST00244	CS	Amyloid A4
1880	EST01634	CS	Axonal glycoprotein TAG-1
1885	EST01639	CS	Histocompatibility antigen modifier 1
. 313	EST00276	CS ·	Lysosomal membrane glycoprotein 1 (LAMP-1)
2017	EST01678	CS	Milk fat globule membrane protein
1567	EST02610	CS	Neural cell adhesion molecule L1
2368	EST01389	CS	Radial spoke protein 3
2089	EST01699	CS	Sperm membrane protein
1277	EST02306	DC	Bib protein
188	EST00256	DC	Enhancer of split
43	EST00371	DC	Maternal G10 protein
1704	EST01580	DC	Myeloid differentiation primary response gene MyD1
. 227	EST00259	DC	Notch/Xotch
952	EST01961	DC	Notch/Xotch
346	EST01828	DC	Orthodentical homeotic protein
1408	EST02442	DC	Seven in absentia
97	EST00289	EM	Aconitase
· 310	EST00377	EM	Fo ATPase beta subunit, mitochondrial
485	EST01466	KP	Calmodulin-dependent protein kinase, type II, beta
. 993	EST02007	KP	Kinase 5 protein
1069	EST02087	KP	Protein kinase C, zeta
1933	EST01650	· KP	Protein phosphatase 2A beta subunit
202	EST00298	KP	Protein-tyrosine phosphatase LRP
1348	EST02378	KP	cAMP-dependent protein kinase inhibitor
2302	EST01779	OG	Discs-large tumor suppressor
2353	EST01806	OG	Prohibitin
1478	EST02515	OG	Rab5
300	EST00232	OG	Transforming protein (dbl)
37	EST00038	OG	ras p21-like small GTP-binding protein (smg GDS)
102	EST00248	OG	rho H12/ ARH12
1834	EST01620	OM	AMP deaminase, brain
691	EST00675	OM	Alcohol dehydrogenase
396	EST01443	OM	CDPdiacylglycerol-serine O-phosphatidyltransferase
2192	EST01257	OM	Diacylglycerol kinase, lymphyocyte
1441	EST02477	OM	Diamine acetyltransferase
2289	EST01325	OM	Fatty acid synthase
1020	EST02034	OM	Glutaminase
2326	EST01791	OM	Inositol-1,4,5-trisphosphate 3-kinase
1427	EST02463	OM	Long-chain-fatty-acid-CoA ligase
2226	EST01744	OM	NAD(P)+ transhydrogenase (B-specific)
1566	EST02609	OM	Neutrophil oxidase factor
1681	EST01573	OM	Nucleoside diphosphate kinase

SEO ID	EST#	Group	Putative Identification
2254	EST01751	OM	Phosphatidylinositol-4,5-bisphosphate phosphodiest
1654	EST01572	OM	Protochlorophyllide reductase
2073	EST01697	OM	Succinate dehydrogenase flavoprotein
2138	EST01715	OM	Succinate dehydrogenase flavoprotein
1771	EST01601	OM	Thiosulfate sulfurtransferase (rhodanese)
2173	EST01724	PI	Lon protease
2297	EST01775	ΡΙ	Prohormone cleavage enzyme
9	EST00376	ΡΙ	Prolyl endopeptidase
1726	EST01588	PI	XPR2 alkaline extracellular protease
1147	EST02169	PP	Tyrosine kinase
2282	EST01764	RT	Lamin B receptor
189	EST00282	RT	trkB
251	EST00370	SC	Actin, other
2146	EST01218	SC	Actin, other
248	EST00271	SC	Actinin, alpha
891	EST01891	SC	Actinin, alpha
1500	EST02538	SC	Actinin, alpha
132	EST00110	SC	Agrin
1852	EST01625	SC	Agrin
2004	EST01676	SC	Cofilin
650	EST00642	SC	Dilute (myosin heavy chain)
2217	EST01738	SC	Gelation factor ABP-280
77	EST00257	SC	Kinesin
· 78	EST00258	SC	Kinesin
2245	EST01748	SC	Kinesin
1468	EST02505	SC	Matrin 3
223	EST00368	SC	Microtubule-associated protein 1B
824	EST01865	SC	Microtubule-associated protein 1B
2032	EST01683	SC	Microtubule-associated protein 1B
506	EST01471	SC	Neuraxin
951	EST01960	SC	Spectrin, beta
1371	EST02402	SC	Talin
653	EST01512	SC	Tubulin, alpha
311	EST00270	SC -	Tubulin, beta
594	EST01490	· SC	Tubulin, beta
757	EST01542	SC	Tubulin, beta
1245	EST02274	SC	Tubulin, beta
1589	EST02634	SC	Tubulin, beta
1701	EST00853	SC	Unc-104
969	EST01982	ST	ADP-ribosylation factor 1
1126	EST02146	ST	Calbindin D28
1910	EST01645	ST	Calmodulin
161	EST00247	ST	MARCKS (myristoylated alanine-rich protein kinase
769	EST00734	ST	MARCKS homolog
1386	EST02418	ST.	MARCKS homolog
1931	EST01041	ST	cAMP-regulated phosphoprotein
1413	EST02447	ST	AMP-specific phosphodiesterase
299	EST00249	ST	smg p25A GDP dissociation inhibitor
•			

SEO ID	EST#	Group	Putative Identification
2092	EST01700	TP	Anion exchanger homolog AE3
1956	EST01663	TP	Ca2+-transporting ATPase 2
1039	EST02055	TP	Calcium channel
1667	EST00825	TP	Gamma-aminobutyric acid transporter
1412	EST02446	TP	Glutamate-aspartate carrier protein
913	EST01913	TT	Clathrin coat assembly protein AP50 homolog
- 1035	EST02051	TT	J1 protein
93	EST00287	TT	Processing enhancing protein
38	EST00374	TT	RNA polymerase II 6th subunit (RPO26)
1715	EST01583	TT	Ribosomal protein L18a
1856	EST01627	TT	Ribosomal protein L1a
1974	EST01667	TT	Ribosomal protein L3
301	EST00300	TT	Ribosomal protein L30
22	EST00301	TT	Ribosomal protein S10
2402	EST01826	TT	Ribosomal protein S10
463	EST01459	TT	Ribosomal protein YL10
- 2121	EST01711	TT	Valine-tRNA ligase
1332	EST02362	TX	GA binding protein, beta subunit
1229	EST02258	TX	KUP protein
1395	EST02429	TX	Nuclear factor 1-like protein (NF1)
· 187 ·	EST00152	TX	Wilm's tumor-related protein
249	EST00275	TX	Zinc Finger Proteins
413	EST01446	TX	Zinc Finger Proteins
469	EST01460	TX	Zinc Finger Proteins
833	EST01560	TX	Zinc Finger Proteins
1230	EST02259	TX	Zinc finger proteins
1496	EST02534	TX	Zinc finger proteins
2324	EST01352	TX.	Zinc Finger Proteins

Group Key: CS: Cell Surface, DC: Developmental Control, EM: Energy Metabolism, KP: Kinases and Phosphatases, OG: Oncogenes, OM: Other Metabolism, PI, Peptidases and Peptidase Inhibitors, RT: Receptors, SC: Structural and Cytoskeletal, ST: Signal Transduction, TP: Transporters, TT: Transcription, Translation, and Subcellular Localization, TX: Transcription Factors.

EXAMPLE 11

<u>cDNA Libraries Generated From Specific Genomic DNA</u> <u>by Exon Expression & Amplification</u>

5

10

15

20

25

30

35

Exon amplification was used to express potential exons from genomic DNA in a recombinant vector that contains some of the signals necessary for splicing. If an exon is present in the proper orientation in the vector, that exon will be spliced in a mammalian cell and will become part of the mRNA of that cell. The exon splice-product can be purified from other mRNA in the cell by conversion of the mRNA to cDNA and selective amplification of the recombinant splice-product cDNAs. Cosmid DNA from human chromosome 19q13.3 was digested with BamHI or BamHI/BglII restriction enzymes. The fragments generated were collected and size specifically cloned into an expression vector (Buckler, et al. Proc. Nat'l. Acad. Sci. USA, 88:4005-4009 After (1991). transfection electroporation of these constructs into COS cells, RNA transcripts were generated using the SV40 early promoter and a polyadenylation signal derived from SV40 both present in the expression vector. When a fragment of genomic DNA contains an entire exon with flanking intron sequence in the sense orientation, the exon should be retained in the mature poly(A) + cytoplasmic RNA. Therefore, the mRNA was used as template for cDNA synthesis using reverse transcriptase and vector-priming. Subsequently, the cDNAs were amplified by vector-priming using PCR. A fraction of this first PCR reamplified using product was internal vector-primers containing terminal cloning sites. These products were endwith T4 DNA polymerase, digested with appropriate restriction enzymes, gel purified and cloned into pBluescript vectors. The constructs were transfected into XL1-Blue competent cells and plated : on LB/Xgal/IPTG/ampicillin plates. White colonies were selected and expanded to prepare DNA templates as described in Example 2.

WO 93/16178

When multiple cosmids or YAC clones were used as the source DNA, a pool of specific expressed exons was obtained as a cDNA library. The EST/cDNAs sequenced from this specific library are disclosed herein as SEQ ID NOS: 2412-2417.

5

10

15

20

25

30

35

EXAMPLE 12

PCR Amplification from Predicted Exons

Computational analyses can be applied to genomic DNA sequences to predict protein coding regions. region prediction program CRM (E. Uberbacher and R. Mural, Proc. Natl. Acad. Sci. USA 88:11261-5 (1991)) finds open reading frames and classifies them according to their probability of being coding regions. These regions are subsequently examined using the GM program (C. Fields and C. Soderlund, Comp. Applic. Biosci. 6: 263, 1990), predicts intron-exon structure. PCR primers are then designed to amplify the predicted exons and used to test human cDNA libraries (for example, fetal brain or placental libraries) for the presence of these putative exons using a PCR assay.

This strategy has been successfully applied in two large scale genomic sequencing projects, the Huntington's locus of human chromosome 4p16.3 (McCombie, et al., submitted) and human chromosome locus 19q13.3 (Martin-Gallardo, et al., submitted). Sequences from eleven predicted exons from chromosome 4 were present in tested cDNA libraries, indicating that this region has at least two and probably three expressed genes. In one case, the method resulted in an amplification product which spanned two predicted exons. (SEQ ID NO: 2411.) When sequenced, this PCR product indicated the presence of the two exons from which the primers were initially chosen, as well as an intervening exon which was also predicted by the CRM program, but not the intervening genomic sequences. In a similar fashion, the presence of the two predicted genes in the chromosome 19

5 .

10

15

20

25

30

35

sequence was confirmed by sequencing PCR products. SEQ ID NO 2410, includes a partial exon of one of these genes.

EXAMPLE 13

Complete Sequence of EST Clone Inserts

There are a number of methods known to those with skill in the art of molecular biology, to obtain sequence information from the cDNAs corresponding to Procedures for these methods are provided in sequences. Basic Methods in Molecular Biology (David et al. supra). One way to acquire more information about the cDNA from which an EST was derived is to sequence the remainder of the cDNA The complete sequence of the inserts of four EST clone. clones (representing SEQ ID NOs 188, 189, 223, and 227) was determined using Exonuclease III deletions. Briefly, EST clones were digested with the restriction enzymes Sall and KpnI or PstI and BamHI (for deletions from the Forward primer and Reverse primer ends of the insert, respectively). The KpnI and PstI enzymes leave 3' sticky ends following digestion, which Exonuclease III is unable to bind. results in unidirectional deletions into the cDNA insert leaving the vector sequence undisturbed. After addition of Exonuclease III to the Forward and Reverse deletion reactions, aliquots of the reaction were removed at defined time intervals and the reaction was stopped to prevent further deletion. S1 nuclease and Klenow DNA polymerase were added to create blunt ended fragments suitable for ligation.

Samples for each time point was purified by electrophoresis through an agarose gel and religated. Two to four representative clones from each time point in each direction were sequenced to give between 200 and 400 base pairs of sequence data. Careful selection of deletion conditions and time points allow a deletion series of approximately 100-200 base pairs difference in length at each consecutive time point. Sequence fragments were reassembled into a redundant contiguous sequence using the INHERIT

software from Applied Biosystems, Inc. (Foster City, CA). In this way, the complete insert from these four cDNA clones was sequenced on both strands to an average redundancy between three and four (each base was sequenced between three and four times, on average). Those complete insert sequences are disclosed herein as SEQ ID 2418, 2419, 2420, and 2421, corresponding to original ESTs with SEQ ID 223, 189, 227, and 188, respectively.

EXAMPLE 14

10

15

20

25

30.

35

5

<u>Determining Reading Frame, Orientation, Coding Regions:</u> <u>ESTs and Complete cDNA Sequences</u>

Once the complete cDNA sequence has been determined in accordance with Example 13, the reading frame, orientation, and coding regions are determined by computer techniques. (The complete coding region is considered to be the largest open reading frame from a methionine to a stop codon.)

Specifically, the CRM program on the GRAIL server is used as explained in Example 9 to determine probable coding regions. This information is supplemented by location of start and stop codons. Where possible, the results of the CRM analysis are validated by comparison of the cDNA sequence to known sequences using database matching, in accordance with Examples 3 and 4. If a match of 50% (or even less) is found in any particular reading frame and orientation, this serves to verify corresponding CRM results. Alternatively, database matches can be used to determine reading frame and orientation without use of the CRM program. Of course, if the cDNA is derived from a directional library, the probable orientation is already known.

EXAMPLE 15

Preparation of PCR Primers and Amplification of DNA

The EST sequences and the corresponding cDNA sequences and genomic sequences may be used, in accordance with the

present invention, to prepare PCR primers for a variety of applications. The PCR primers are preferably at least 15 bases, and more preferably at least 18 bases in length. procedure of Example 5 is repeated using the desired EST, or using the corresponding cDNA or genomic DNA sequence from Example 13. It is preferred that the primer pairs have approximately the same G/C ratio, so that temperatures are approximately the same. When screening cDNA, introns are of no concern; however, when screening genomic DNA, primers should be selected to avoid reading across introns, which usually are too large to amplify. PCR primers and amplified DNA of this Example find use in the Examples that follow.

15

20

35

5

EXAMPLE 16

Forensic Matching by DNA Sequencing

In one exemplary method, DNA samples are isolated from forensic specimens of, for example, hair, semen, blood or skin cells by conventional methods. A panel of PCR primers derived from a number of the sequences of Example 1, 2, 11, 12 and/or 13 is then utilized in accordance with Example 12 to obtain DNA of approximately 100-200 bases in length from the forensic specimen. Corresponding sequences are obtained from a suspect. Each of these identification DNAs is then sequenced, and a simple database comparison determines the differences, if any, between the sequences from the suspect and those from the sample. Statistically significant differences between the suspect's DNA sequences and those from the sample conclusively prove a lack of identity. lack of identity can be proven, for example, with only one sequence. Identity, on the other hand, demonstrated with a large number of sequences, all matching. Preferably, a minimum of 50 statistically identical sequences

WO 93/16178 PCT/US93/01294

-79-

of 100 bases in length are used to prove identity between the suspect and the sample.

EXAMPLE 17

5

10

15

Positive Identification by DNA Sequencing

The technique outlined in the previous example may also be used on a larger scale to provide a unique fingerprinttype identification of any individual. In this technique. primers are prepared from a large number of sequences from Examples 1, 2, 11, 12 and/or 13. Preferably, 20 to 50 different primers are used. These primers are used to obtain a corresponding number of PCR-generated DNA segments from the individual in question in accordance with Example 15. of these DNA segments is sequenced, using the methods set forth in Example 1. The database of sequences generated through this procedure uniquely identifies the individual from whom the sequences were obtained. The same panel of primers may then be used at any later time to absolutely correlate tissue or other biological specimen with that individual.

EXAMPLE 18

25

30

35

20

Southern Blot Forensic Identification

The procedure of Example 17 is repeated to obtain a panel of from 10 to 2000 amplified sequences from an individual and a specimen. This PCR-generated DNA is then digested with one or a combination of, preferably, four base specific restriction enzymes. Such enzymes are commercially available and known to those of skill in the art. After digestion, the resultant gene fragments are size separated in multiple duplicate wells on an agarose gel and transferred to nitrocellulose using Southern blotting techniques well known to those with skill in the art. For a review of Southern

blotting see Davis et al. (<u>Basic Methods in Molecular</u> <u>Biology</u>, 1986, Elsevier Press. pp 62-65).

A panel of ESTs or complete cDNA sequences from Examples 1, 2, and/or 13, or fragments thereof of at least 15 bases, are radioactively or colorimetrically labeled using endlabeled oligonucleotides derived from the ESTs. nick translated sequences or the like using methods known in the art and hybridized to the Southern blot using techniques known in the art (Davis et al., supra). Preferably, at least 5 to 10 of these labeled probes are used, and more preferably at least about 20 or 30 are used to provide a unique pattern. The resultant bands appearing from the hybridization of a large sample of ESTs will be a unique identifier. Since the restriction enzyme cleavage will be different for every individual, the band pattern on the Southern blot will also be unique. Increasing the number of EST probes will provide statistically higher level of confidence in the identification since there will be an increased number of sets of bands used for identification.

20

30

35

5

10

15

EXAMPLE 19

Dot Blot Identification Procedure

25 Another technique for identifying individuals using the sequences disclosed herein utilizes a dot blot hybridization

technique.

Genomic DNA is isolated from nuclei of subject to be identified. Oligonucleotide probes of approximately 30 bp in length were synthesized that correspond to sequences from the The probes are used to hybridize to the genomic DNA through conditions known to those in the The oligonucleotides labelled P³² are end with using polynucleotide kinase (Pharmacia). Dot Blots are created by spotting about 50 ng cDNA of at least 10, preferably at least 50 sequences corresponding to a variety of the Sequence ID

WO 93/16178 PCT/US93/01294

-81-

NOs provided in Table 7 onto nitrocellulose or the like using a vacuum dot blot manifold (BioRad, Richmond California). The nitrocellulose filter containing the EST clone sequences is baked or UV linked to the filter, prehybridized and hybridized with labeled probe using techniques known in the art (Davis et al. <u>supra</u>). The 32P labeled DNA fragments are sequentially hybridized with successively conditions to detect minimal differences between the 30 bp sequence and the DNA. Tetramethylammonium chloride is useful for identifying clones containing small numbers of nucleotide (Wood et al., mismatches Proc. Natl. Acad. 82(6):1585-1588 (1985) which is hereby incorporated by A unique pattern of dots distinguishes one reference. individual from another individuals.

15

20

25

30

35

10

5

EXAMPLE 20

Alternative "Fingerprint" Identification Technique

EST sequences and the corresponding complete cDNA sequences can be used to create a unique fingerprint for an individual. Thus pools of EST sequences can be used in forensics, paternity suits or the like to differentiate one individual from another.

Entire EST sequences can be used; similarly oligonucleotides can be prepared from EST sequences. In this example, 20-mer oligonucleotides are prepared from 200 EST sequences using commercially available oligonucleotide services such as Oligos Etc., Wilsonville, OR. Patient cell samples are processed for DNA using techniques well known to those with skill in the art. The nucleic acid is digested with restriction enzymes EcoRI and XbaI. digestion, samples are applied to wells for electrophoresis. The procedure, as known in the art, may be modified to accommodate polyacrylamide electrophoresis, however in this example, samples containing 5 ug of DNA are loaded into wells

5

10

15

20

25

30

35

and separated on 0.8% agarose gels. The gels are transferred using Southern blotting techniques onto nitrocellulose.

10 ng of each of the oligos are pooled and end-labeled with P³². The nitrocellulose is prehybridized with blocking solution and hybridized with the labeled probes. Following hybridization and washing, the nitrocellulose filter is exposed to X-Omat AR X-ray film. The resulting hybridization pattern will be unique for each individual.

It is additionally contemplated within this example that the representative number of EST sequences can be varied for additional accuracy or clarity.

EXAMPLE 21

Identification of genes associated with hereditary diseases

This example illustrates an approach useful for the association of EST sequences with particular phenotypic characteristics. In this example, a particular EST is used as a test probe to associate that EST with a particular phenotypic characteristic.

An EST clone corresponding to EST01643, (SEQ ID NO 1894) maps to a gene rich region of chromosome 6. EST clone HHCMH89, from which EST01643 was derived, was mapped to chromosome 6p21 by Dr. Julie Korenberg of UCLA/Cedar Sinai Hospital using FISH. A search of Mendelian Inheritance in Man (supra) revealed 6p21 to be a very gene rich region containing several known genes and several diseases for which genes have not been identified. The cDNA encoded by EST clone HHCMH89 thus becomes an immediate candidate for each of these genetic diseases.

Cells from patients with these diseases are isolated and expanded in culture. PCR primers from the EST sequences are used to screen genomic DNA and RNA or cDNA from the patients. ESTs that are not amplified in the patients can be positively associated with a particular disease by further are lysic.

EXAMPLE 22

Identification of a gene associated with Angelman's disease

5

10

15

. 20

25

30

35

Angelman's disease (AD) is characterized by deletions on the long arm of chromosome 15 (15q11q13) (Williams et al. Am. J. Med. Genet. 32:339-345 (1989) hereby incorporated by The symptoms of the disease include developmental delay, seizures, inappropriate laughter and ataxic movements. These symptoms suggest that the disorder neurologic deficiency. This prophetic example illustrates how ESTs, preferably obtained from a cDNA library from human brain, may be used in identifying the defective gene or genes associated with Angelman's Disease. example is based on analogous work with genomic DNA, rather than cDNA and ESTs, in identifying the genetic defect associated with Angelman's Disease.) This example also illustrates how EST sequences may generally be used for identifying gene sequences associated with an inherited disease that is mapped to a chromosome location.

ESTs are screened using techniques described in Example 5 and Example 7 to identify those ESTs that localize to the long arm of chromosome 15 and preferably localize to chromosome 15 bands 15q11q13 from normal patients. ESTs that bind to the long arm of chromosome 15 are hybridized to chromosome 15 from AD patients. These studies preferrably performed using either fluorescence in situ hybridization or using somatic cell hybrids that contain fragments from the long arm of chromosome 15 from AD Those chromosome 15-specific ESTs that do not map patients. to chromosome 15 from AD patients are useful as markers for Angelman's Disease and can be incorporated into diagnostics for genetic screening. These ESTs are associated with chromosome deletions in present Angelman s disease. Identification of the gene associated with these AD negative ESTs and an analysis of the polypeptides encoded by the genes

10

15

20

25

from normal patients is essential for providing gene or other therapies for AD patients.

Genetic diseases are not always accompanied by gene deletions. Therefore, it is also important to use the ESTs that bind to bands 15q11q13 from AD patients as tools to identify the polymorphisms present within the disease population. Restriction fragment length polymorphism (RFLP) analysis can be performed on patient cells from AD disease or from somatic cell hybrids created using the long arm of chromosome 15. For a review of RFLP techniques see Donis-Keller et al. (Cell 51:319-337 (1987) hereby incorporated by DNA is isolated from the somatic cell lines or reference). from cells from AD patients. The DNA is digested with one or more restriction enzymes according to techniques of Donis-Keller et al. The resulting fragments are separated by gel electrophoresis, denatured, transferred to nitrocellulose and hybridized with the selected radio-labeled ESTs that localize to the region of interest. The autoradiographic pattern is compared both to a number of AD patients and to normal Common patterns of EST hybridization in AD patients that are not present in normal patients indicates that the genes associated with these ESTs are candidate genes affected by AD.

cDNA libraries are prepared from the somatic cell hybrids from AD patients. Libraries are prepared using Lambda Zap II Library Kits (Stratagene, La Jolla, California) or other commercially available library kits. The ESTs of interest are used as probes to identify those bacterial colonies carrying genes corresponding to the EST probes. Positive clones are sequenced and the sequences are compared to homologous gene sequences derived from normal patients.

Alterations, including deletions and substitutions, within gene sequences, associated with bands 15q11q13, are thus positively identified and associated with AD disease. Wagstaff et al. were able to identify deletions and substitutions in sequences encoding the GABA, receptor

10

15

20

25

.30

35

protein subunit from patients with Angelman's disease (Am. J. Hum. Genet. 49:330-337, (1991)). It is likely that other genes will additionally be associated with the disease.

5 EXAMPLE 23

Preparation and Use of Antisense Oligonucleotides

Antisense RNA molecules are known to be useful for regulating translation within the cell. Antisense RNA molecules can be produced from EST sequences or from the corresponding gene sequences. These antisense molecules can be used as diagnostic probes to determine whether or not a particular gene is expressed in a cell. Similarly, the antisense molecules can be used as a therapeutic to regulate gene expression once the EST is associated with a particular disease (see Example 22).

The antisense molecules are obtained from a nucleotide sequence by reversing the orientation of the coding region with regard to the promoter. Thus, the antisense RNA is complementary to the corresponding mRNA. For a review of antisense design see Green et al., Ann. Rev. Biochem. 55:569-597 (1986), which is hereby incorporated by reference. The antisense sequences can contain modified sugar phosphate backbones to increase stability and make them less sensitive to RNase activity. Examples of the modifications are described by Rossi et al., Pharmacol. Ther. 50(2):245-254, (1991).

Antisense molecules are introduced into cells that express the gene corresponding to the EST of interest in culture. In a preferred application of this invention, the polypeptide encoded by the gene is first identified, so that the effectiveness of antisense inhibition on translation can be monitored using techniques that include but are not limited to antibody-mediated tests such as RIAs and ELISA, functional assays, or radiolabelling. The antisense molecule is introduced into the cells by diffusion or by transfection

procedures known in the art. The molecules are introduced onto cell samples at a number of different concentrations preferably between $1\times10^{-10}M$ to $1\times10^{-4}M$. Once the minimum concentration that can adequately control translation is identified, the optimized dose is translated into a dosage suitable for use in vivo. For example, an inhibiting concentration in culture of 1×10^{-7} translates into a dose of approximately 0.6 mg/kg bodyweight. Levels of oligonucleotide approaching 100 mg/kg bodyweight or higher may be possible after testing the toxicity of the oligonucleotide in laboratory animals.

The antisense can be introduced into the body as a bare or naked oligonucleotide, oligonucleotide encapsulated in lipid, oligonucleotide sequence encapsidated by viral protein, or as oligonucleotide contained in an expression vector such as those described in Example 25. The antisense oligonucleotide is preferably introduced into the vertebrate by injection. It is additionally contemplated that cells from the vertebrate are removed, treated with the antisense oligonucleotide, and reintroduced into the vertebrate. It is further contemplated that the antisense oligonucleotide sequence is incorporated into a ribozyme sequence to enable the antisense to bind and cleave its target. For technical applications of ribozyme and antisense oligonucleotides see Rossi et al.

EXAMPLE 24

Preparation and use of Triple Helix Probes

30

35

25

10

15

20

Triple helix oligonucleotides are used to inhibit transcription from a genome. They are particularly useful for studying alterations in cell activity as it is associated with a particular gene. The EST sequences or complete sequences of the present invention or, more preferably, a portion of those sequences, can be used to inhibit gene

5

10

15

20

25

30

35

expression in individuals having diseases associated with a Similarly, a portion of the EST particular gene. corresponding gene sequence can be used to study the effect of inhibiting transcription of a particular gene within a Traditionally, homopurine sequences were considered the most useful. However, homopyrimidine sequences can also inhibit gene expression. Thus, both types of sequences from either the EST or from the gene corresponding to the EST are contemplated within the scope of this invention. Homopyrimidine oligonucleotides bind to the major groove at homopurine: homopyrimidine sequences. As an example, 10-mer to 20-mer homopyrimidine sequences from the ESTs can be used to inhibit expression from homopurine sequences. SEQ ID NOs such as 282, 888, 719, 670, 994, 240, 873 and 761 contain homopyrimidine 15-mers. Moreover the natural (beta) anomers of the oligonucleotide units can be replaced with alpha anomers to render the oligonucleotide more resistant to nucleases. Further, an intercalating agent such as ethidium bromide, or the like, can be attached to the 3' end of the alpha oligonucleotide to stabilize the triple helix. information on the generation of oligonucleotides suitable for triple helix formation see Griffin et al. **245**:967-971 (1989), which is hereby incorporated by this reference).

The oligonucleotides may be prepared an oligonucleotide synthesizer or they may be purchased commercially from a company specializing in oligonucleotide synthesis. The sequences are introduced into cells in culture using techniques known in the art that include but are not limited to calcium phosphate precipitation, DEAE-Dextran, electroporation, liposomemediated transfection or native uptake. Treated cells are monitored for altered cell function. These cell functions are predicted based upon the homologies of the gene. corresponding to the EST from which the oligonucleotide was derived, with known genes sequences that have been associated WO 93/16178

with a particular function. The cell functions can also be predicted based on the presence of abnormal physiologies within cells derived from individuals with a particular inherited disease, particularly when the EST is associated with the disease using techniques described in Example 22.

EXAMPLE 25

Gene expression from DNA Sequences Corresponding to ESTs

10

15

20

25

30

35

A gene sequence of the present invention coding for all or part of a human gene product is introduced into an expression vector using conventional technology. (Techniques to transfer cloned sequences into expression vectors that direct protein translation in mammalian, yeast, insect or bacterial expression systems are well known in the art.) Commercially available vectors and expression systems are available from a variety of suppliers including Stratagene (La Jolla, California), Promega (Madison, Wisconsin), and Invitrogen (San Diego, California). If desired, to enhance expression and facilitate proper protein folding, the codon context and codon pairing of the sequence may be optimized for the particular expression organism, as explained by Hatfield, et al., U.S. Patent No. 5,082,767, incorporated herein by this reference.

The following is provided as one exemplary method to generate polypeptide from cloned cDNA sequences. The cDNA from the EST of interest is sequenced to identify the methionine initiation codon for the gene and the poly A sequence. If the cDNA lacks a poly A sequence, this sequence can be added to the construct by, for example, splicing out the Poly A sequence from pSG5 (Stratagene) using BglI and SalI restriction endonuclease enzymes and incorporating it into the mammalian expression vector pXT1 (Stratagene). pXT1 contains the LTRs and a portion of the gag gene from Moloney Murine Leukemia Virus. The position of the LTRs in the construct allow efficient stable transfection. The vector

WO 93/16178 PCT/US93/01294

includes the Herpes Simplex Thymidine Kinase promoter and the selectable neomycin gene. The cDNA is obtained by PCR from the bacterial vector oligonucleotide using primers complementary to the cDNA and containing restriction endonuclease sequences for Pst I incorporated into the 5'primer and BglII at the 5' end of the corresponding cDNA 3' primer, taking care to ensure that the cDNA is positioned inframe with the poly A sequence. The purified fragment obtained from the resulting PCR reaction is digested with PstI, blunt ended with an exonuclease, digested with Bgl II, purified and ligated to pXT1, now containing a poly A sequence and digested BglII.

5

10

15

20

25

30

35

The ligated product is transfected into mouse NIH 3T3 cells using Lipofectin (Life Technologies, Inc., Grand Island, New York) under conditions outlined in the product specification. Positive transfectants are selected after growing the transfected cells in 600ug/ml G418 (Sigma, St. Louis, Missouri). The protein is preferrably released into the supernatant. However if the protein has membrane binding domains, the protein may additionally be retained within the cell or expression may be restricted to the cell surface.

Since it may be necessary to purify and locate the transfected product, synthetic 15-mer peptides synthesized from the predicted cDNA sequence are injected into mice to generate antibody to the polypeptide encoded by the cDNA.

If antibody production is not possible, the cDNA sequence is additionally incorporated into eukaryotic expression vectors and expressed as a chimeric with, for example, β -globin. Antibody to β -globin is used to purify the chimeric. Corresponding protease cleavage sites engineered between the β -globin gene and the cDNA are then used to separate the two polypeptide fragments from one another after translation. One useful expression vector for generating β -globin chimerics is pSG5 (Stratagene). This vector encodes rabbit β -globin. Intron II of the rabbit β -globin gene facilitates splicing of the expressed transcript,

the polyadenylation signal incorporated into the construct increases the level of expression. These techniques as described are well known to those skilled in the art of molecular biology. Standard methods are published in methods texts such as Davis et al. and many of the methods are available from the technical assistance representatives from Stratagene, Life Technologies, Inc., or Promega. Polypeptide may additionally be produced from either construct using in vitro translation systems such as In vitro $Express^{TM}$ Translation Kit (Stratagene).

Example 26

Production of an Antibody to a Human Protein

15

20

25

30

35

10

Substantially pure protein or polypeptide is isolated from the transfected or transformed cells as described in Example 25. Concentration of protein in the final preparation is adjusted, for example, by concentration on an Amicon filter device, to the level of a few micrograms/ml. Monoclonal or polyclonal antibody to the protein can then be prepared as follows:

A. Monoclonal Antibody Production by Hybridoma Fusion

Monoclonal antibody to epitopes of any of the peptides identified and isolated as described can be prepared from murine hybridomas according to the classical method of Kohler, G. and Milstein, C., Nature 256:495 (1975) or derivative methods thereof. Briefly, a mouse is repetitively inoculated with a few micrograms of the selected protein over a period of a few weeks. The mouse is then sacrificed, and the antibody producing cells of the spleen isolated. The spleen cells are fused by means of polyethylene glycol with mouse myeloma cells, and the excess unfused cells destroyed by growth of the system on selective media comprising aminopterin (HAT media). The successfully fused cells are diluted and aliquote of the dilution placed in wells of a

microtiter plate where growth of the culture is continued. Antibody-producing clones are identified by detection of antibody in the supernatant fluid of the wells by immunoassay procedures, such as Elisa, as originally described by Engvall, E., Meth. Enzymol. 70:419 (1980), and derivative methods thereof. Selected positive clones can be expanded and their monoclonal antibody product harvested for use. Detailed procedures for monoclonal antibody production are described in Davis, L. et al. Basic Methods in Molecular Biology Elsevier, New York. Section 21-2.

B. Polyclonal Antibody Production by Immunization

5

10

15

20

25

30

35

Polyclonal antiserum containing antibodies heterogenous epitopes of a single protein can be prepared by immunizing suitable animals with the expressed protein. described above, which can be unmodified or modified to enhance immunogenicity. Effective polyclonal antibody production is affected by many factors related both to the antigen and the host species. For example, small molecules tend to be less immunogenic than other and may require the use of carriers and adjuvant. Also, host animals vary in response to site of inoculations and dose, with both inadequate or excessive doses of antigen resulting in low titer antisera. Small doses (ng level) of antigen administered at multiple intradermal sites appears to be most reliable. An effective immunization protocol for rabbits can be found in Vaitukaitis, J. et al. J. Clin. Endocrinol. Metab. 33:988-991 (1971).

Booster injections can be given at regular intervals, and antiserum harvested when antibody titer thereof, as determined semi-quantitatively, for example, by double immunodiffusion in agar against known concentrations of the antigen, begins to fall. See, for example, Ouchterlony, O. et al., Chap. 19 in: Handbook of Experimental Immunology D. Wier (ed) Blackwell (1973). Plateau concentration of antibody is usually in the range of 0.1 to 0.2 mg/ml of serum (about 12 μ M). Affinity of the antisera for the antigen is

determined by preparing competitive binding curves, as described, for example, by Fisher, D., Chap. 42 in: Manual of Clinical Immunology, 2d Ed. (Rose and Friedman, eds.) Amer. Soc. For Microbiol., Washington, D.C. (1980).

Antibody preparations prepared according to either protocol are useful in quantitative immunoassays which determine concentrations of antigen-bearing substances in biological samples; they are also used semi-quantitatively or qualitatively to identify the presence of antigen in a biological sample.

EXAMPLE 27

Identification of Tissue Types or Cell Species by Means of Labeled Tissue Specific Antibodies

15

20

10

Identification of specific tissues is accomplished by the visualization of tissue specific antigens by means of antibody preparations according to Example 26 which are conjugated, directly or indirectly to a detectable marker. Selected labeled antibody species bind to their specific antigen binding partner in tissue sections, cell suspensions, or in extracts of soluble proteins from a tissue sample to provide a pattern for qualitative or semi-qualitative interpretation.

25

30

Antisera for these procedures must have a potency exceeding that of the native preparation, and for that reason, antibodies are concentrated to a mg/ml level by isolation of the gamma globulin fraction, for example, by ion-exchange chromatography orby ammonium sulfate fractionation. Also, to provide the most specific antisera, unwanted antibodies, for example to common proteins, must be removed from the gamma globulin fraction, for example by means of insoluble immunoabsorbents, before the antibodies labeled with the marker. Either monoclonal heterologous antisera is suitable for either procedure.

35

A. Immunohistochemical Techniques

5

10

15

20

25

30

35

Purified, high-titer antibodies, prepared as described above, are conjugated to a detectable marker, as described, for example, by Fudenberg, H., Chap. 26 in: Basic & Clinical Immunology, 3rd Ed. Lange, Los Altos, California (1980) or Rose, N. et al., Chap. 12 in: Methods in Immunodiagnosis, 2d Ed. John Wiley & Sons, New York (1980).

A fluorescent marker, either fluorescein or rhodamine, is preferred, but antibodies can also be labeled with an enzyme that supports a color producing reaction with a substrate, such as horseradish peroxidase. Markers can be added to tissue-bound antibody in a second step, as described below. Alternatively, the specific antitissue antibodies can be labeled with ferritin or other electron dense particles, and localization of the ferritin coupled antigen-antibody complexes achieved by means of an electron microscope. In yet another approach, the antibodies are radiolabeled, with, for example ¹²⁵I, and detected by overlaying the antibody treated preparation with photographic emulsion.

Preparations to carry out the procedures can comprise monoclonal or polyclonal antibodies to a single gene copy or protein, identified as specific to a tissue type, for example, brain tissue, or antibody preparations to several antigenically distinct tissue specific antigens can be used in panels, independently or in mixtures, as required.

Tissue sections and cell suspensions are prepared for immunohistochemical examination according common histological techniques. Multiple cryostat sections (about 4 μ m, unfixed) of the unknown tissue and known control, are mounted and each slide covered with different dilutions of the antibody preparation. Sections of known and unknown tissues should also be treated with preparations to provide a positive control, a negative control, for example, preimmune sera, and a control for non-specific staining, for example, buffer.

Treated sections are incubated in a humid chamber for 30 min at room temperature, rinsed, then washed in buffer for 30-45 min. Excess fluid is blotted away, and the marker developed.

5

10

15

20

25

30

If the tissue specific antibody was not labeled in the first incubation, it can be labeled at this time in a second antibody-antibody reaction, for example, by adding fluorescein- or enzyme-conjugated antibody against the immunoglobulin class of the antiserum-producing species, for example, fluorescein labeled antibody to mouse IgG. Such labeled sera are commercially available.

The antigen found in the tissues by the above procedure can be quantified by measuring the intensity of color or fluorescence on the tissue section, and calibrating that signal using appropriate standards.

B. Identification of Tissue Specific Soluble Proteins

The visualization of tissue specific proteins and identification of unknown tissues from that procedure is carried out using the labeled antibody reagents and detection strategy as described for immunohistochemistry; however the sample is prepared according to an electrophoretic technique to distribute the proteins extracted from the tissue in an orderly array on the basis of molecular weight for detection.

A tissue sample is homogenized using a Virtis apparatus; cell suspensions are disrupted by Dounce homogenization or osmotic lysis, using detergents in either case as required to disrupt cell membranes, as is the practice in the art. Insoluble cell components such as nuclei, microsomes, and membrane fragments are removed by ultracentrifugation, and the soluble protein-containing fraction concentrated if necessary and reserved for analysis.

A sample of the soluble protein solution is resolved into individual protein species by conventional SDS polyacrylamide electrophoresis as described, for example, by Davis, L. et al., Section 19-2 in: Basic Methods in Molecular Biology (P. Leder, ed), Elsevier, New York (1986), using a

range of amounts of polyacrylamide in a set of gels to resolve the entire molecular weight range of proteins to be detected in the sample. A size marker is run in parallel for purposes of estimating molecular weights of the constituent proteins. Sample size for analysis is a convenient volume of from 5-50 μ l, and containing from about 1 to 100 μ g protein. An aliquot of each of the resolved proteins is transferred by blotting to a nitrocellulose filter paper, a process that maintains the pattern of resolution. Multiple copies are prepared. The procedure, known as Western Blot Analysis; is well described in Davis, L. et al., (above) Section 19-3. One set of nitrocellulose blots is stained with Coomassie Blue dye to visualize the entire set of proteins for comparison with the antibody bound proteins. The remaining nitrocellulose filters are then incubated with a solution of one or more specific antisera to tissue specific proteins prepared as described in Example 26. In this procedure, as in procedure A above, appropriate positive and negative sample and reagent controls are run.

5

10

15

20

25

30

35

In either procedure A or B, a detectable label can be attached to the primary tissue antigen-primary antibody complex according to various strategies and permutations thereof. In a straightforward approach, the primary specific antibody can be labeled; alternatively, the unlabeled complex can be bound by a labeled secondary anti-IgG antibody. In other approaches, either the primary or secondary antibody is conjugated to a biotin molecule, which can, in a subsequent step, bind an avidin conjugated marker. According to yet another strategy, enzyme labeled or radioactive protein A, which has the property of binding to any IgG, is bound in a final step to either the primary or secondary antibody.

The visualization of tissue specific antigen binding at levels above those seen in control tissues to one or more tissue specific antibodies, prepared from the gene sequences identified from EST sequences, can identify tissues of unknown origin, for example, forensic samples, or

differentiated tumor tissue that has metastasized to foreign bodily sites.

The entire contents of all references cit d above are hereby incorporated by reference.

While the present invention has been described in some detail for purposes of clarity and understanding, one skill d in the art will appreciate that various changes in form and detail can be made without departing from the true scope of the invention.

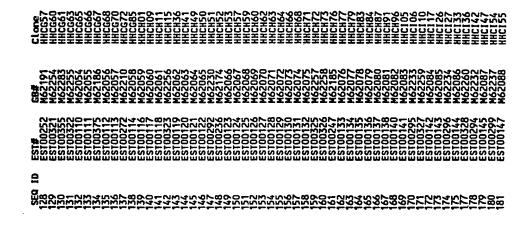
10

VII. Correlation of EST and Clone Identifiers

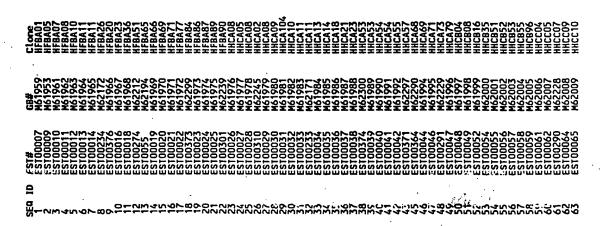
The EST sequences of the present invention are identified herein by SEQ ID NO, and are identified in the GenBank database by a different number, are identified in the inventors' lab (and upcoming publications) by EST number, and clones have been submitted to the American Type Culture Collection (Rockville, Maryland USA) under clone names. Table 12 cross references those different numbers for the ESTs from cDNA, SEQ ID NOS 1-2409.

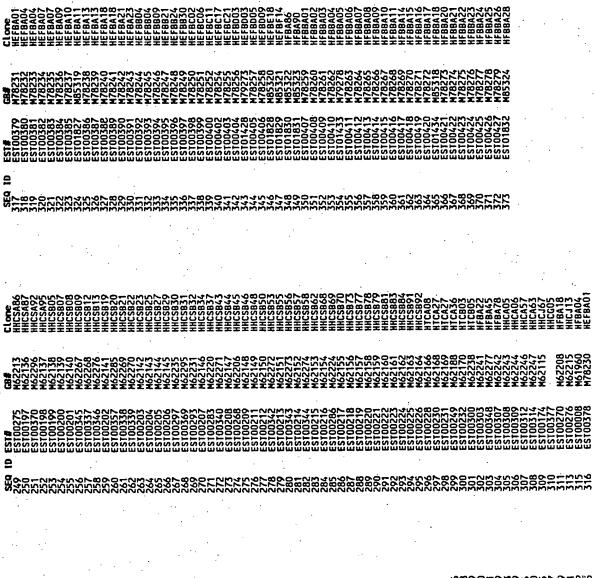
20

Certain Sequence ID NOS are excluded from some claims based on their homology to known non-human sequences (S e Table 2).

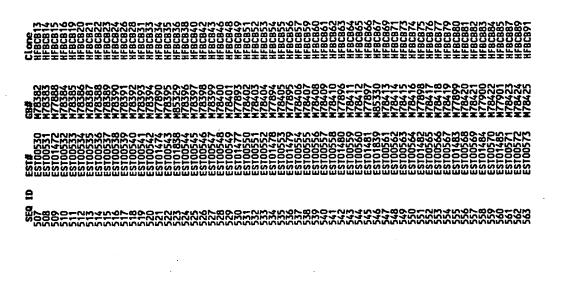


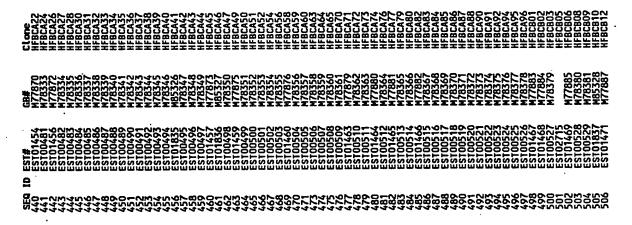


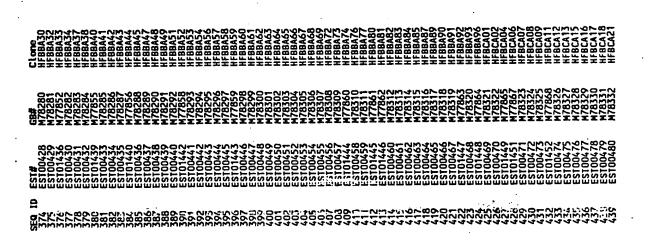


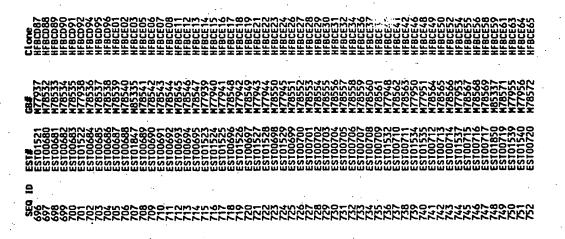


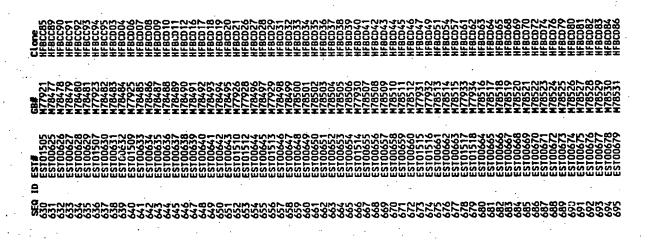
| Colored Box |

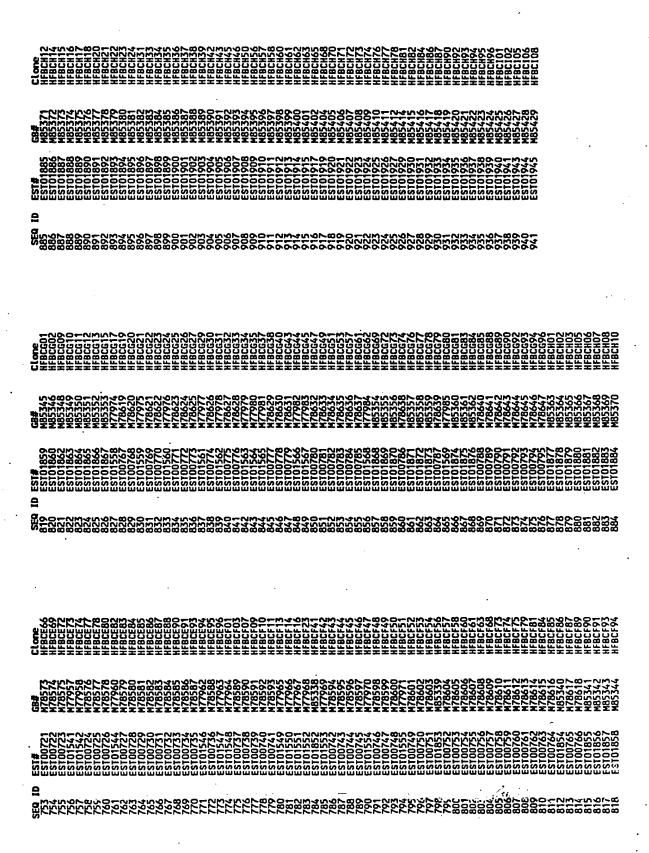


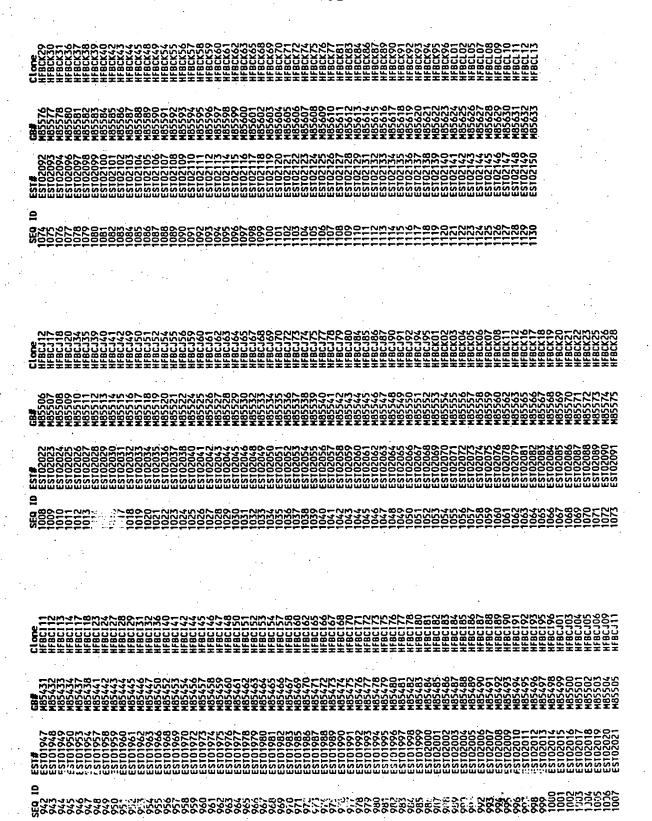


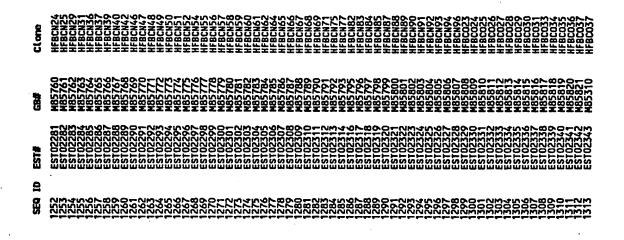


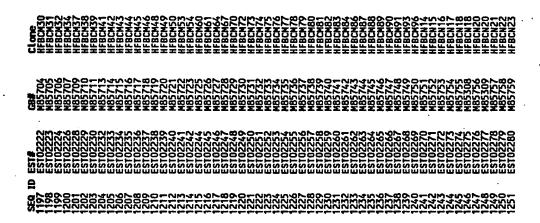


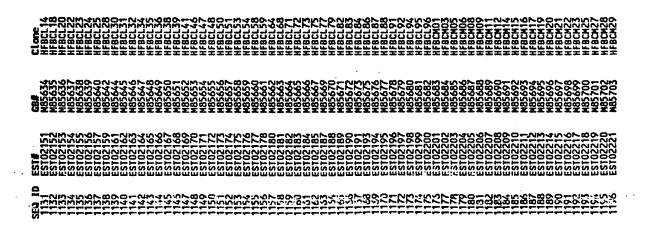


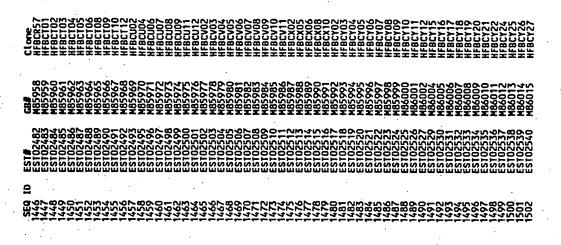


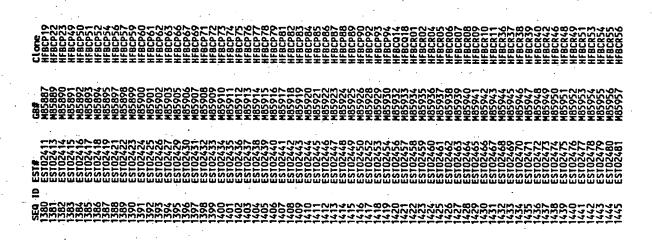


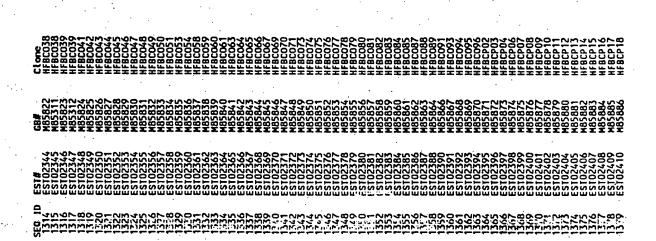


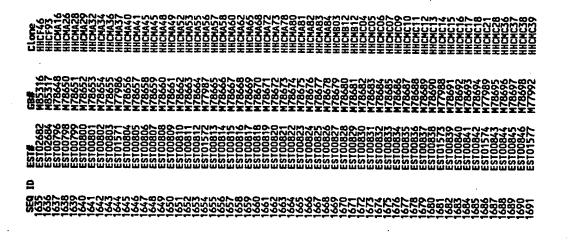


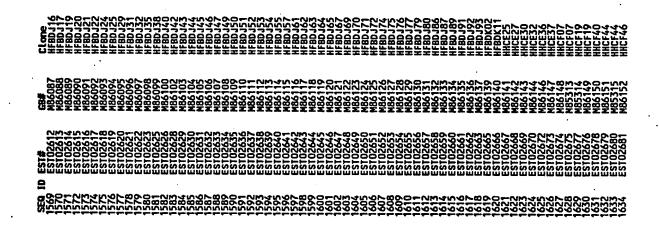


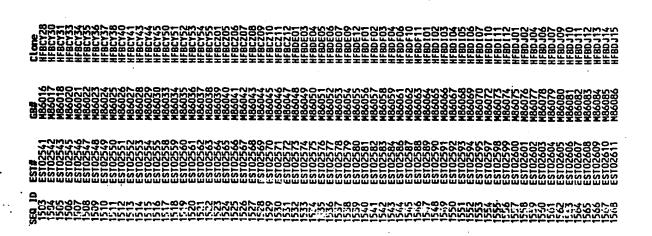


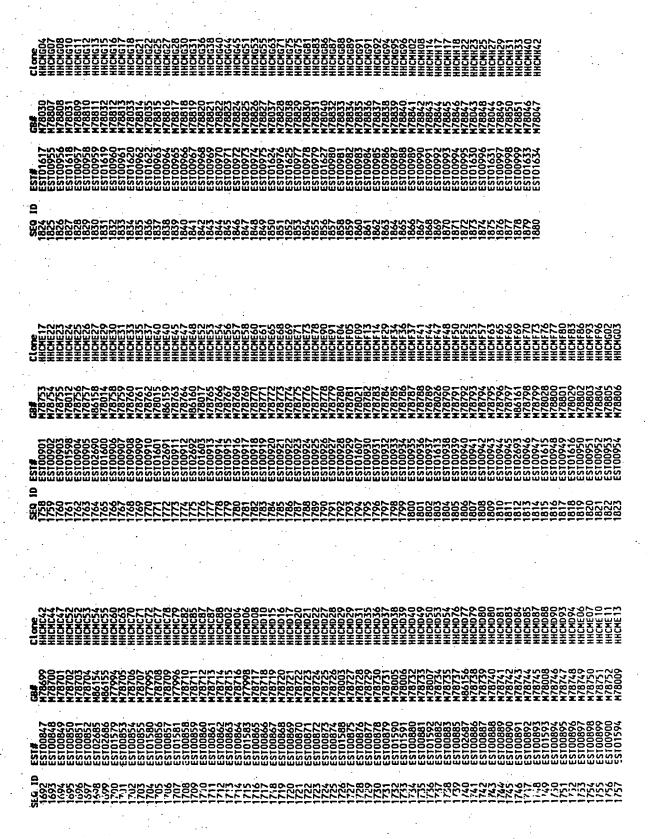


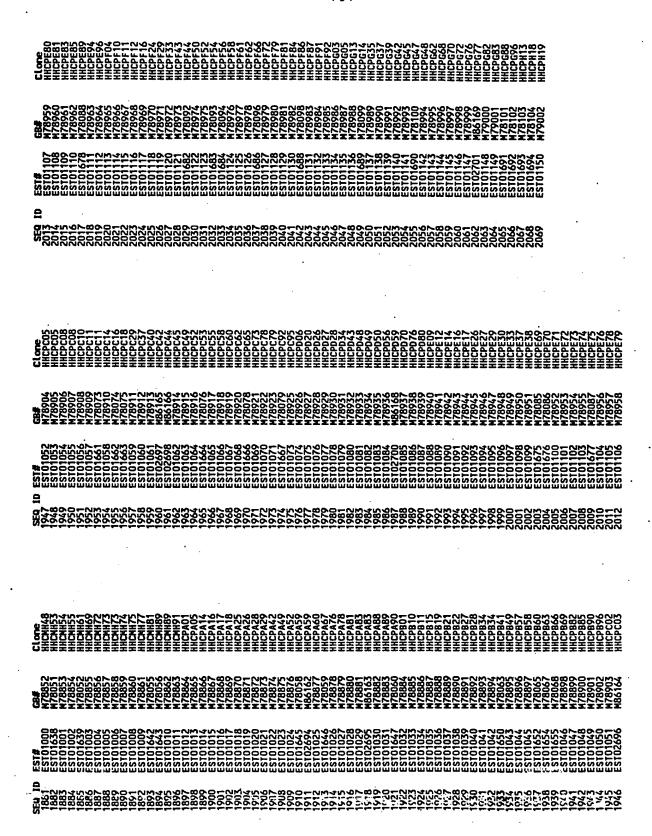


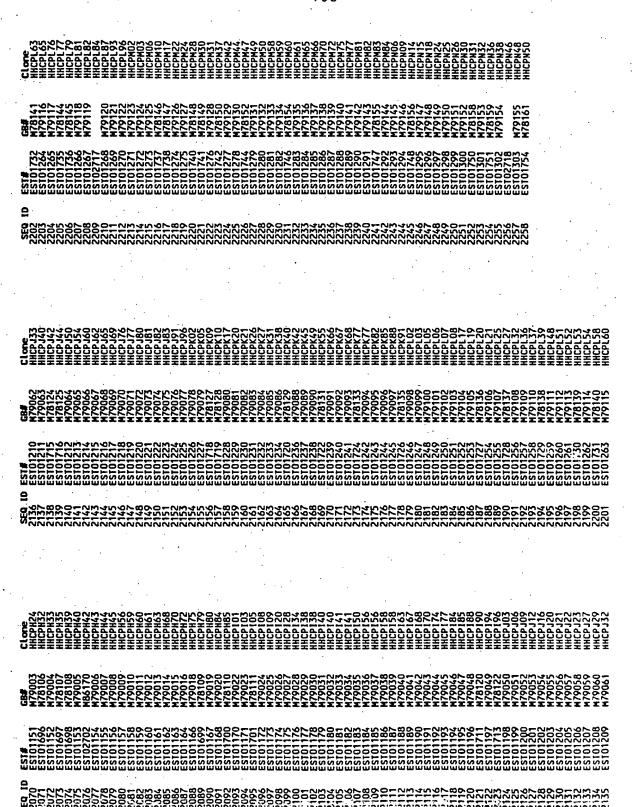


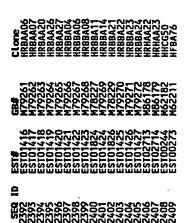


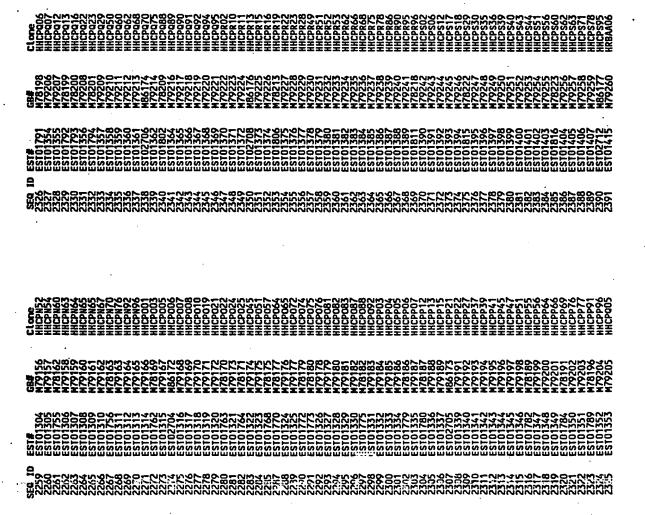












NOTE REGARDING SEQUENCE LISTINGS: The listings of SEQ ID NOS: 1-2421 are in numerical order. However, an occasional number (for example, SEQ ID NO: 44) is not found in this list. In all, 9 SEQ ID NOS are not used. Nevertheless, the convention "1-2421" is used, for example, to refer to all the SEQ ID NOS in the following list, while "1-315" is used, for example, to refer to all the listed sequences falling between SEQ ID NO 1 and SEQ ID NO 315.

.

SEQUENCE LISTING

- (1) GENERAL INFORMATION:
 - (i) APPLICANT: Venter, J. Craig

Adams, Mark D.

Moreno, Ruben F.

- (ii) TITLE OF INVENTION: Sequences Characteristic of Human Gene Transcription Product
- (iii) NUMBER OF SEQUENCES: 2412 (1-2421, with 9 SEQ ID NOS unused.)
 - (iv) CORRESPONDENCE ADDRESS:
 - (A) ADDRESSEE: Knobbe, Martens, Olson, and Bear
 - (B) STREET: 620 Newport Center Dr. Sixteenth Floor
 - (C) CITY: Newport Beach
 - (D) STATE: CA
 - (E) COUNTRY: USA
 - (F) ZIP: 92660
 - (v) COMPUTER READABLE FORM:
 - (A) MEDIUM TYPE: Floppy disk
 - (B) COMPUTER: IBM PC compatible
 - (C) OPERATING SYSTEM: PC-DOS/MS-DOS
 - (D) SOFTWARE: PatentIn Release #1.0, Version #1.25
 - (vi) CURRENT APPLICATION DATA:
 - (A) APPLICATION NUMBER: 07/837,195
 - (B) FILING DATE: 12-FEB-1992
- (vii) PRIOR APPLICATION DATA:
 - (A) APPLICATION NUMBER: US 07/716,831
 - (B) FILING DATE: 20-JUN-1991
- (viii) ATTORNEY/AGENT INFORMATION:
 - (A) NAME: Israelsen, Ned A.
 - (B) REGISTRATION NUMBER: 29,655
 - (C) REFERENCE/DOCKET NUMBER: NIH004.004CP1
 - (ix) TELECOMMUNICATION INFORMATION:
 - (A) TELEPHONE: 619-235-8550
 - (B) TELEFAX: 619-235-0176

SEO ID NO:1: (Length of Sequence = 362 Nucleotides)

CITCCCTTT GITCCCCTCA GIGICCCTT TAATTGCTTC CCTCCATTTT CCTTAGCAGC ATCCTAGTEG ATGGTCTGGGG
TTATCAGAGG AGCAAAAACA TITAAGTGTC AAATAATGCT CATTGTCTCC CTGGGATTTC TAAACAGAAA AAATGAAGAA

AGAGGCAGAG AAGAGCTTCA CAAGGTGTGT GCCAGCTCTG CATCATTTCC AGCTGCTCAA CCACCATTTC TCCCATTTTA
GGTCCCCAAA AGTAGGAGGT GGGGCCTCAC AGAGCTGCTG TGGGCTTTGG GTATCAAAAG CTGCAGCCAC CATATGGGGC
ACTCCTGGCT GGTGTACAGG GTGGGCATTG CCCAGGTCTT TT

SEO ID NO:2: (Length of Sequence = 214 Nucleotides)

GITTINCTIT TITCTIAGCI TCATTICTCI TAAAAAACAA GGAACAAGAA AACATIGCAC CAGCGITCIA AGCCICAAAC
AAAANACAAA ACAAATCCCC CIGCGAAGAA CAATAAACII TACATCICTI IGGCAACAAT AACIITAAAAT CACCCAACIII
CCATICGCIC CAACCACAGC AGIIAGIIAG TIACAAAAAT ATICCNIGIG CIGC

<u>SEO ID NO:3:</u> (Length of Sequence = 344 Nucleotides)

ATTAATAGGA AAGATGATTG TATAGATGGT GGGCTATTAA CTCAGATCAG GATGAGAATC GGGAGTGCCT TTACATGTGT GGTACCCAAA TGGGTGGTTG GATATAAGAG TAACAAAAGG ACTGAAAGGG TTAAAAAAGA AAGAAAAAAA AAAAACTCCC TGGTTGGGAG GGTGTTAAGT ATCGAGTGTT TTTCCAAACC ATTCCTCCTC TGCTCACCTA CCCCTAGGTG ATTAAAGGAG ATTACCTTTTA AAAAAGAAAG AATTGGCTCA AAGGTACTGT AAATTCTAGG ATTATATACC TTTATATAGG TTCATTCCCT GATCCCTGTA TTATCAAGGC ACAG

SEO ID NO:4: (Length of Sequence = 352 Nucleotides)

GACCCGGTAA CCGAGGCGGC AAGGAGGCCA GGTAGTCCCG GCACCTCTCA CTCTGCAGAG ACCAGCGGCT TCGTGGGAGG CCTGTGGGTC ACACGTAGGG GCTAGAGCCA GCCTGCATCC TGCCCACCGG GCTCCACTTG GAGATCAGCA GGAGGGCCAG TGTGGGACCC CTGCTGCCAC CTCTCCTGGG CCTGTKTCCT TTCTGGAAAT TAAGAAGGTG TGCTCCAGAG CCAAGAGGAG CAATAAGAAA CCTCGTGTGC CAGCTTCTTA AGGGTKGCAG TGCAAGACCC CA

SEO ID NO:5: (Length of Sequence = 562 Nucleotides)

ATACCCTIAC ATATATATIC ACAGAAAATC ATATIGCATA TACTICITICT CCACATCATA AAAATGGGTG TIGGGCTCTC
TAGGACACAA GGGAAGCAGG CCAAATTICT CATATITICA GGAATAAACT GAGTGCCCCG AAGGTGTAAT AGGAACCTTT
TACTAACCTC ATCTGACTTC ATCCTCACAC CAGCATTITG TGTGTAAGGA AACTGGCCGA GAGTGGTTAA GAAATATATC
CAAAGACGTA TAGTTCCAAA TGGAACACGG ATCTTTTAT TTAAATTCCA ATCATCTTTC CATTATATCA GCCAATGATG
GAGCAGAAAG CTGGTCCAGG CAATCCCAGA ATAGATCTTT CTAGGCACCC GTTCAGTGTG AGGAGGGGGA AGTGGCCTTG
CCAAAGGCCC AGTGAGCTCA ATTAGGGTTA ACGCTGCTTC TTAGCCTACC CCAGGGGNCA CCGCACTTAG GTTGTTTTGT
GCCCAGCTTT GGCAGGAAGC ATTCCTCCTT TCAAAGATTN NAGCCTTGCG GTCATATATC GGGTGTAATA GGGTTCTTTT
TT

SEO ID NO:6: (Length of Sequence = 359 Nucleotides)

ACATGITCIC CCICTITCAA TITTAGCAGI AATGIGATCC TCAAAAATGC ATTAATACTA GITGAAGTAA ATAAACGGAA GAGCICCAAA AIGCCIGCAT TAAATGCAIT TITCCACACT AAIGCCAATC AICCAAAGCT ATTITCAACA AGICAGGTAT TCAAAGCTAT TCACACCACT TGAAAGAGTA AITACCATTI ACIGAAGCAC TTATCTGICC TACACTGATG GGAGTAAATG CITCICATAG GITATCTCAT GITACATTATG CCACITINAC TTAAAATGAT CACAATINAG TGCTATAGGT TITTGGGTTA AITGITTTCCC NGGGGGAGIT GITAAAAACA TGGCATTIC

SEO ID NO:7: (Length of Sequence = 218 Nucleotides)

AACITGCAAC ATAAATACTA GAAAAAGAGA AAATATCATC AAAATACAAA TAACTGTTAG AAATCATTGC TCAAAAGAAR AACCTGGCAA TGCATGATTA CGAAATGCAA AAGAMGATAC AGTTGCTCTC TGTATATGCG CTTTCCACAT CCACAGATTC AAACAACTGT GGATAAAAAA GGATTTTTCA ATGCCATTAA ACAVCAATGC AACAGTAA

SEO ID NO:8: (Length of Sequence = 345 Nucleotides)

CTACAATAGA AGGCAAACTA TGICCCICCT TIGCICAGAA ACITITAATA TCIKCCIATT TCCCCATGIA AAAGCCAATC
CTCAACCACA GIGIAGAAGG GCIATCCAIT TCIAGCIACA CATCTCCTCA GICACTGCCC CCAGCCCCAG TACTTGGGGA
CTTTGCCCTT GCAGTICCCT GIGCCAGCAA ACICTTCCTC CAGATGICCA CATGACTCAC CCINCTCCTT CAGGGGTCTT
CTCAAATGIC ACITTACCAG AGGIGGCTTC CCIGACCATC CIGIATAAAT AGCATCACCC TACCTCCTAT CTCTCTCTCT
AATGICTCAG GAATTCGATA TCAAG

SEO ID NO:9: (Length of Sequence = 189 Nucleotides)

GIGAACAGAC TAAGGCCTTT NIGGAGGCCC AGAATAAGAT TACTGTGCCA TITCTTGAGC AGTGTCCCAT CAGAGGTTTA
TACAAAGAGA GAATGACTGA ACTATATGAT TATCCCANGT ATAGTTGCCA CTTCAAGAAA GGAGAACGGT GTTTTTATTT
TTACAAATACA GGNTTINAGA ACCACCGG

SEO ID NO:10: (Length of Sequence = 267 Nucleotides)

CTCCCTTCGC CACCTGCTGG ACGCGAGGG CTACTACGAT GCCATGGGTG TCCTGRTTT TTATTTCTCA GACAGGACTG-CTCTGTAINT GTCTTTGGAT TCTACGTAGA TTTATATTTG TAAAATATTA CATTTGTCAT GACCAGAAGA AATGTCATTA TCGTAAAATT TAGATTCTGG NGTCTATATA TGNAAGNAAT ACTAACTACT AACTGTTATA ACAWCAAAAT GTGGGNTGTA TATCTACARG CCNGAGCCGA CTTGTCA

SEO ID NO:11: (Length of Sequence = 247 Nucleotides)

CTCATAAAGC CAGGGGATA AAAWIGGTAG TITCATGITA TCTACAAGRC TAAGKICAAA ATTCCATGCA TGGGCGGRTA AAAGACCCAT NATGGKCCIM ACTGTACTTA CICCCCATTT ATTAGCATTC ATTCTGGTCA CCAGGTCTAG TTCCTCTGCT TAGCGAATCT CGCTTGTCTT CAAGATGTCA TTCAAATGTC ACATTTTGTG GGAAGCCTTG CCTTTTTTGA CACGGTCTCC CTGCCAC

SEO ID NO:12: (Length of Sequence = 280 Nucleotides)

AAGGCGAGAG GCTTCTGGAG AAACCCACCC CACCAACGTC TTGATCTTGG ACTTTTAVCC TCCAGAGCTA TGAGAAAACA
AVTTTCTGTV VATVGVGGCC ACTCAGCCTG TGGATACTGG CAGCCCTAGC AAACTCATAC ACACATACAT TTTAAACTCG
GTTTAATCCT GTGRCCATTC ACTTATGGTT CAGTTTTTAA ATAGTCCTAG TCTTATGVCC ACTGTTAAAG TTCACCAGGA
CATAGGSCAT TGGGGAAAGG GGCCTGTAAC TCTTGGATTA

SEO ID NO:13: (Length of Sequence = 339 Nucleotides)

VCIVICIVCE AACTICATIC AGATATIGAC ICIGGIGAIG GGAACATIAA ATACATICIC ICAGGGGAAG GAGCIGGAAC
CATTITIVIR ATTGATGACA AATCAGGGAA CATTCATGCC ACCAAGACGI IGGATCGAGA AGAGAGAGCC CAGTACACGI
TGATGGCICA GGCGGIGGAC AGGGACACCA ATCGGCCACT GGAGCCACCG ICGGAATICA ITKICAAGGK CCAGGACAIT
AATGACAGIC CICCGGAGGI ITCCIGCACG AGACCIATCA IGCCAACIGI GCCSIGIARA GGICCAATKI IGGGIGSIGI
ACGGIAGIGG GGAGGCCIG

SEO ID NO:14: (Length of Sequence = 342 Nucleotides)

GGGYGCAAAG TAGCAGATTC TAGTAAAGGA CCAGATGAGG CAAAAATTAA GGCACTCTTG GAAAGAACAG GCTACACACT
TGATGTGACC ACTGGACAGA GGAAGTATGG AGGACCACCT CCAGATTCCG TITATYCAGG TCAGCAGCCT TCTGTTGGCA
CTGAGATATT TGTGGGAAAG ATCCCAAGAG ATCTATTTTG AGGATGAACT TGTTCCATTA TTTGAGAAAG CTTGGACCTA
TATGGGATCC TTCGTCTAAT GATGGATCCA CTCACTGGTC TCAATAGAGG TTAATGCGTT TGTCACTTTT TTGTACAAAA
GGAGCARGCT CAAGGAGGGC TG

SEO ID NO:15: (Length of Sequence = 354 Nucleotides)

ATGITGATGC TGAAATTVAA GATCCACCAA TTCCAGAAAA ACCATGGAAG GITCATGIGA AATGGATITT GGACACIGAT
ATTITCAATG AATGGATGAA TGAGGAGGAT TATRAGGIGG ATGAAAATAG GAAGCCIGIR AGIITYCGIC AGCGGATITC
AACCAAGAAT GAAGAGCCAG TCAGAAGTCC AGAAAGAAGA GATAGAAAAG CATCASCTAA TGCTCGAAAG AGGAAACATT
CGCCTTCGCC TCCCCCTCCG ACACCAACAG AWTCACCGGA AGAAGAGTCG GAAGAAAGGC CAAGCTAGCC TTTTATGGGG
AAGCCGCAAG AAGTCCAGAA AGAGGGWWGG TTGA

SEO ID NO:16: (Length of Sequence = 348 Mucleotides)

CAGGCAAGTT TCTTCCAGGA TGAGAAATCA GTGGAAAGTG AGGGCCAGCC AACAGCCACC ACCAACCACC CAACACGCGA GCGAGACCAT CTTAAAAGAG CCCCAGCCAA GCTGACCATG GGTCTGACCC CAAACTGAAG AAATGCCCAG CCCAGCCAAA CCCCAAATTGC TAACTTGTAT TATAAGCAAG TACAATGGTC CTTACCTTAA GCCACTAAGT TTTGGGATGC TTTGTTACAC AGCTATAGAT AAGCTGATAC AGGGAATGTC AGAWTCCATG ATGAGAGACC GAGCCTTTCA KTCTGTCAGA GGYACCTTVG GTTGGCAAAA CTTCAAAAAG AGGGACCT

SEO ID NO:17: (Length of Sequence = 415 Nucleotides)

AGCAYGGGCT GGGGGGCCGG GAGITAGGGC TGGGGCTIGI TITTACGCTCT GCCCCCCACA CCCCCTCCTC TTCCGTCCTG
ATTAAGCCCA AGGGITGGIG GACITAACTT TCAGCCCATC TCTAAGGGIT TCACAGACTG GATCITTCTA AACTITATTG
GGTACCTGCT TCCCCTTTTC CCTGGIAGIT TTCATCTACA AAAAGTCAAA ACCTGATCGA AATAGAAATA AGATCATCAA
ATTGGACCAT TCTCTTAGCG TTCGAGIGIG CCGGCCAGAC TGGCATTCAG TACACGCTGA GATCCAACCA CATCACACTG
GCCTCAGGTC ACCAACTCGC CACTCAGGGC ACAAGGCCTG CCCTTGTGGT CACAAGGCTT TCCTTAATGT CGTCGGTGCC
CAGGTGAACC ACAAG

SEO ID NO:18: (Length of Sequence = 356 Nucleotides)

GIATGIAGTA TICIATACIT AACCATCIGI GICCCAATTA AGCTAAACAT GATTCATTCI GATGCCAACC
CCCATCCATC ATGCCATGGA TCGCTCTAGA CTTCTTCCCT TGTAACCTCC CACTCAAACA GIGAGAAACC TTTGCCCAGT
ATGTTTTGGA GIAACCTCAC TGGGAGTTG CAGTCCCACT AGATGAATGC CAACCCATTT GITCATTTAA AAGGACTTTT
GGAACCATAG AGCAATGGCT GGGCTGGGTC TVGCACGTTC ATCTTGACTG AAACAATTGG CCATGAAGGC ACTTGCCCAAG
GAAACTCTAG GGGCCACAAG GGTCCTGGGT GCTTGC

SEO ID NO:19: (Length of Sequence = 339 Nucleotides)

CATGCTTCCA TITITITIAG TITTAAACCA CCAAACCAAT ATTITYCCTT TAAATTTAA TCITATAATA TAGAAATCTT
ATGTAAATGA AATTITGTCA TGTTTCAAAT AAAGAGAACT GAAGTAGAAA ATAGAAATGC CAGTAAACAA CATAATGTTT
AAATTACAAC TTACATTAGG GGTTTGGGGG VATGCTAATT ATATATTGAG AATATACATT AGAACTCTTC AAAATGGGCT
CTTCTAATGA GGTCACTACT GAACATAATT GTTCCCTCTT CTGTTAAATA GAATAGGTTT AAATGACTAG TCCAAATGGA
ATTATTGCCT TCTKGTTAA

...............................

SEO ID NO:20: (Length of Sequence = 437 Nucleotides)

AGAACAAGGG AACTCAGCAG CCCCTCCCTT CCCATCAGCT GTTCCTGAGA GATGCAATAT AGTAGTCATC GACATCATCC
TTATCAACAG CATCATCACT CAGACAGTGG TGAAAGTCTT TCTTCACAAG GAAAAACAAA GATAAAGAAA TACATGAGCA
TTAATCAGAA ATTTTCAAAG CTTGGATTCT AATGATATGC ATTATCATTA GACATTCAAA TGCTATACAT CTTCTGATGA
AGCCTCCTTG ACAGCAGCTA CACTTATTTC ACATTAGAAT GCCTAGAGAA ATCCTGACTG CCCAGCTTGG TCATGGGACC
TTCCCCCACTC TCCTCTTGGA GGAATGAAAA GATGTGGCGG CTTTCTACTT TTGCTACTGA GCTGGGGTAT ATGGCTAGGT
CCACTTTCTA AGGGGCTTGG AAGGGTTATT CCCATCTG

SEO ID NO:21: (Length of Sequence = 385 Nucleotides)

GITTGATTIG CITTTITTI AGAGITTIAC ATCAGIGITI TICAGGAATA TIGGICITIC ATTITCITIT CITGGAATAT
TITCTAGITI TACTITGICA GAGIAAATIC TGGCITCACA GAATIATTIG TAGICCICC TGICTIGGIT TATTCATGCI
GCTATAACAA AATACCACAG ACAAGGIGGI AATAAATAAC ACAAATTIAT TITTCCCAGI TCIGGAGGCI AGGAGITCAA
GAAGCIGGCA AGITCAATGI CIGGIGAGAC CCATTCCTIC ATAGGIGGCA CCATCIAGGG GICCITACAT GRCAAAGAGA
TGGAAGGGCC AAAAAGATGG TGACCIATTG TGAGGCCTIT TITAAAGGGC CITVAAATCC CAGIC

SEO ID NO:22: (Length of Sequence = 374 Nucleotides)

ACCITCATGE TCATGAAGGC CATGCAGTCT CTCAAGTCCC GAGGCTACGT GAAGGAACAG TITGCCTGGA GACATTTCTA
CTGGTACCTT ACCAATGAGG GTATCCAGTA TCTCCGTGAT TACCTTCATC TGCCCCCGGA GATTGTGCCT GCCACCCTAC
GCCGTAGCCG TCCAGAGACT GGCAGGCCTC GGCCTAAAGG TCTGGGAGGG TGAGCGACCT GCGAGACTCA CAAGAGGGGA
AGCTGACAAG AGATACCTAC AAGACGGGAG TRCCTGTGCC ACCTGGTGCC GACAAGAAAG CCGAGGCTTG GGTCTGGGTC
AGCAACCGAA TTCCAGTTTA GAGGCGGATT TVGGTCGTKG ACGGTGTCAG CCAC

SEO ID NO:23: (Length of Sequence = 322 Nucleotides)

CAAAACGIGA TCACCACAGC TCCGITCCIG CAGIGACACT TAACATACIC AGCATCITCA TGAATTCIGA ATAATTTACT
GATCGIAAAAGIA TCAATITCAG GIGAGCAGIT TIAAATCAGA AAATAGTCAA TAGITAATCA TGACTCITCA
GGGIATTTCC TTCACGICCT CIGAAGAGIT TCCCAGAACA TTCITGIGAA AAGGAATGCC TCCCAACAAT GGAGAGCAAC
AATAGCAACA GGCATCIGAA TCAGCCIGGC CICTGAAAAC AGACCANAGA GGAGITTATC TGITTCITCC AGIGGAGGAA
GG

SEQ ID NO:24: (Length of Sequence = 113 Nucleotides)

CCTGAAATCG GAGTCTTTTG GACTGACTCC AAATTCAATG GGTGGCACAG GCAGCACGGA GTCCACGTGA ATCTCCACCC CGTTAACAGG CGGGACGACA GCCCCTTGCA GCC

SEO ID NO:25: (Length of Sequence = 399 Nucleotides)

GEARAGART AAGGAAAAAC AAGACAAAAT CTACTTCATG GCTGGGTCCA GCAGAAAAGA GCAGACGCTG GCCTCAGACA
CAGACAGCAG TCTTGATGCC TCGACGGGAC CCCTTGAAGG CTGTCGATGA TAGGITAGAA ATAGCAAACC TGTCAGCATT
GAAGGAACTC TCACCTCCGT GGGCCTGAAA TGCTTGGGAG TTGATGGAAC CAAATAGAAA AACTCCATGT TCTGCATGTA
AGAAACACAA TGCCTTGCCC TACTCAGACC TGATAGGAT GCCTGCTTAG ATGATAAAAT GAGGCAGAAT ATGTCTTGAA
GAAAAAAANTT GCAAGCCACA CTTCINGAGA TTTTGTTCAA GATCCATTTC AGGGTGAGCA GTTAGAGTAG GTTGAATTT

.....

SEC ID NO:26: (Length of Sequence = 350 Mucleotides)

GATTGGTATA CGGCCAACAA TGGATTGATA GCCTTAATAT AGAAATAGTT CCAGCAGGCC AGATGCAGTG GCTCAATTCT
GTAAACCCAG TGCTCTGCAC AGCTAGGAAG GAAGATCACT TGGGCCCAGG AGTTCAAGGC TCCAGTGAGC CATGATCACG
CCACTKCCTC CAGCCTGGGT GACAGAGTNA GGCCCTGTCT CTAAAAAATG AAATAGCTCC ATCAAGTCAA TAATTAAAAG
TTCAACAGCC CAACAGANCA AAAATTGTAA ATGANCACAA ATTAGAAAAT GTACAAATTA AATATTAATG ACCCATAACC
CTATAAGGGA AAGTTTAACC TCTCTAGTAT TTTTT

SEO ID NO:27: (Length of Sequence = 322 Mucleotides)

AAAACGTGAT CACCACAGCT CCGTTCCTGC AGTGACACTT AACATACTCA GCATCTTCAT GAATTCTGAA TAATTTACTG
ATCGTAAAAGT CTAAAAGTAT CAATTTCAGG TGAGCAGTTT TAAATCAGAA AATAGTCAAT AGTTAATCAT GACTCTTCAG
GGTATTTCCT TCACGTCCTC TGAAGAGTTT CCCAGAACAT TCTTGTGAAA AGGAATGCCT CCCAACAATG GAGGAGCAAC
AATAGCAACA GGCATCTGAA TCAGCCTGGG CTCTGAAAAC AGACCAAAGA GGNGTTTTC TGCTTTCTTC CAGTGAGGAA
GG

SEO ID NO:28: (Length of Sequence = 287 Nucleotides)

TATTITIATI AAAGGACCAC CCTGGCTGIM GTGAGATGAA TGGATTCAAA CAGGGCAAGA GTGGATACAG MGAGATAAGT
TAGGAAGCTG GTATAGAAAT CTGGATGAGA TATGGTGGCT TGGATGATAC TAGCAGTGAG TATGGGAAGT AGGTGGATTA
CTTTACACTT TTTTAGATCA GTCKATTCTT GATGTCTTGA AGACAAATTA ATCTCATATA TAACTCTAAA CAACATATTT
ATATTTCATG TAAATAAGGA TAATGCTGAC CAAATATTAG CACCTTT

SEO ID NO:29: (Length of Sequence = 282 Nucleotides)

CAGGGCAGGG AAGCCIGGAA GCAAAGGAGG ACCIGGCICC TGACTCICAG AGAGGATAGG CIGGGATCCC TGGGGCAGGC CIGITCCITG GCIGGCCAAT TIAGICTITC AATTGICTAA GGGCICTCCA TIGCCIGCCC TIGCCICITT CIAGCCIGIT ATTICIAGGC TCCICIGAAT AAATCICAGG TITCCIACIG TCATGCCITT AGITCAAAAA TGAGAATCIG CCCTACAGIG CIGGCCICCT TCCGGCCIGA AAGCCAGCAC CITKCGACCC GG

SEO ID NO:30: (Length of Sequence = 345 Nucleotides)

GAAGCIGGIG AATACATITC AAGACACAAC AIGGCACCIG IGICIAGCIC TAIGGIACAA CAIGGIACTA IGACACATAT
AAIGGGIIGC CAGAIGGGGA AGGCAGCIIC ICIGCAACIG AGCIGAGAIC ICAAAATAGA CAAIGICAAG AIGGAAIGAG
AAGGGAAAAA CAGCAIGIGI AGACAGGIAG IGACAAAAGG CIAAITAAGG ACIGAAAGAA ACCAGIGGCC AACAAGGGAA
ICIACGGGIG AIAAAGAIAA GACGGIGAGA GAGAIAAGGC IAGAITGIAI AAGGCIIGAC AGACCAIAGC AAGAIAAGCA
AGGACCIGIG ICCIGIIAAC CAITI

SEO ID NO:31: (Length of Sequence = 343 Nucleotides)

ATAAAATTGG TCTGGGTACC CTAAGGTGTT TGCKTTGATA GAAAATTGAC ACCCCAAACT AAGTGTTCTA CTTAGCTTCT ACAATAGTTA TTCCTAGACC TTAGATTAGT CATTACATTT TTATTTAAGG TACTATGTTA CTTTCATGAC TACAAAATGA GGCACTCGTA CAAAACAGGA ATGAAAACAT ACATATACTG TCTTGTCTTT ATGTCGTATT AATGCCAAAG ATATTGTCAG GGATTATTTT AAAGAAGCCC TTACTCATGA TGGCTATTTT TAAAAATGGC ACAGGACAGT AACAGGCTGA AAAGAAACAC CTGGTTTGAG GGGCCAAATT AAG

SEO ID NO:32: (Length of Sequence = 153 Nucleotides)

ACAGGATGGT CAGGACAAGC CACCTCTGGT AAAGTGACAT TTGAGANGAC CCCTGAAGGN GGGGGGTTGA GTCATGTGGA CATCTTGAGG AAGAGTTTAC TGGCACAGGG AACTGCAAGG KCAAAGTCCC CAAGTACTAG GGCTGGGGGC AGT

SEQ ID NO:33: (Length of Sequence = 257 Nucleotides)

TCAGTCAGCT TATCGCAGGT GCAGCCAAAC ACAAAGCTTC AGGACAAATT GTACAAACTT TACAATGTGG GATTTAAATT
TAAAATATGA TACATAAAAA TCTACACAAA ACTGATAAAA ATCAAGCACA GNTACCAGGA TTGAAACTTA TAATAATCCA
TGTGTGAAAG GGAGTCTTGT TTCCTTTCAA GTGCTTTTAT TCTGCTATGG AACAGTCAAA ATGGAAGNTG TAAAGCTTTG
TGGTTAGTTT AAATTAT

SEO ID NO:34: (Length of Sequence = 307 Nucleotides)

CTCCCACCCA TATCTAATCC AACAAGTCCA GCTGCCTCTC TCTNAAMAAT ACCNARGATC AGGCCCCTTC TCAGCACCCC CACAGCTGCT GCCCCAAAAGG AAGCCACGGTC ATCTCTCACG GAGATTGTKC AGCAGCCACT GCCTCCTTGT CACCTTCGCC TGTGGTCATT CTCCCCACAT GGCCAGGGAA TGCGTCCTGT TAAAGTCTGC TAGGTCACGG TCCTTCCTAC TCAAAATGCT CCCYTGGCTC CCACTGCCCC CAGAGTAAAA AGCCCAGACC TTCAAATGAC ACAAAGGCCT ACAACGA

SEO ID NO:35: (Length of Sequence = 266 Nucleotides)

TCCACAGGTC ATCAGATRCC TGCINGATAA TATATAAACA GIAAAAACAA CITTCACITC TICCTATINI AATCGIGIGC CATGGATCIG ATCIGIACCA TGACCCIACA TAAGGCIGGA TGGACCICAG GCIGAGGGCC CAATGIATGI KIGGCIGIGG GIGIGGITGG GAGIGIGICT GCKGAGIAAG AACACGNITT TCAAGATICT AAAGCICAAT TMAAGIGGCA CATTAATRAT AAACTCAGAT CIGNICAAAA GICCGG

SEQ ID NO:36: (Length of Sequence = 388 Nucleotides)

CAGCITTGGA AAGACTITGA CCTCTGAACA AAAAGCCAGA AGGCTGCTTA AAGAAATAGT AAGGGITTCA CTTGCCCTGG
ATAGTCACAA ATCTAGGAGT ACTGGTTCAC TGCCTTGGGT TACCAGGTAT CAGCTCTTC ACAATCTCTC CTCTTCCCAT
GCTTCCCCTT AAAGTCCAGT TGACAAATGA AAAAGAAAAA AAGGCCTTGA TTTATAGTAT TGCCAAACAA CCTCATAAGA
ATGGGTAAAA TTACATACAC ACATACATAG AGAAGGGAGG TAATGCTGTG AATCTACTTG AGCTGGATTG CATGCTCCCT
AGGGACCACG GTGCCCAACC TGTAATTTTA TTTCTAACTT TTATAAATAT ACTCCTTTTT CACGGATG

SEO ID NO:37: (Length of Sequence = 342 Nucleotides)

GAATGICIAC ACAAGGAAGI ACAGGATTIG GCITTICIAG ATGICATATC CAAACITCGC AGICATGAGA ACAAAAGIGI
TGCCCAGCAG GCCICTCICA CAGAGCAGAG ACITACIGIG GAAAGCIGAG AACIGCCCGA TACACGGCAT CATCCCATCT
CTAATTICCC CICTIGICCIC CATCCAGCGG CITCTICCGC TICATTCTCT ACCATACCAC TIGIGCATGC ATGIRATGIT
CTAATACCAA TIGAAGAACC GCIGIAGGIA CCICCCTAAT AAGGATTICT AAACCTATAG TIAGIGIGAT CATGACITTIG
GTCAAAGGCA AGTYTCCCAC CC

SEO ID NO:38: (Length of Sequence = 355 Nucleotides)

GATGACITGG AGAATGCCGA AGAGGAAGGC CAGGAGAATG TCGAGATCCT CCCCTCTGGG GAGCGACCGC AGCCAACCAG
AAGCGAATCA CCACACCATA CATGACCAAG TACGAGCGAG CCCGCGTGCT GGGCACCCGA GCGCTCCAGA TTGCGATGTG
TGCCCCTGTG ATGGTGGAGC TGGAGGGGGA GACAGATCCT CTGCTCATTG CCATGAAGGA ACTCAAGGCC CGAAAGATCC
CCATCATCAT TCGCCGTTAC CTGCCAGATG GGAGCTATGA AGACTGGGGG GGTKGACGAG CTCATCATCA CCGACTTGAG
CTGGAGTCAT CTTTCCTGXC CTTTGCCCCA TGCCC

SEQ ID NO:39: (Length of Sequence = 303 Nucleotides)

GCCAAAAACA NYTCTGAACC CGITTTGGGA AATAATGGGA TTCCTTGATC ACGGGACAAC GAATCACCCT GAAGTTTTTC
TCCAGTTTAC TCAGTCACAT AAGCCACCAG AGGCTAACCA CACTGACAAC AAAAGCAAGT CCCAGGATTC CGGGGGCTAA
TACCATGCTA GGCATTACTT GGGAAGTTAT GAGTTGGTAT ACATCTGTGA ATTTGGTGGG AGGAGAAAAC TAACAGTAAA
TTTATCAAAG CCAGTGGTAC GTTCAGCGTT ATAAAAATTA CAAGGATCTG CTTCTCGGCG ACT

<u>SEO ID NO:40:</u> (Length of Sequence = 178 Nucleotides)

GGTGTCGGGG GCTAGAGATA CACATGCCAG INCIATACAT TTCTCAGCAC TGTGCTGTCG ATTCACAGCA GTTCAATTGT TCATGCGATA TAAGCCAGTC ATGTGGCCCA AGTTATTCTG TCGGCTGTGT TCTCTGCAGG AATCTGATGC AAGAAGGCCT GAAGGATGCA TGGCTTTT

<u>SEQ ID NO:41:</u> (Length of Sequence = 322 Nucleotides)

TECCTITCTT TAGAAATTTA GGGCAGIGIG ATGCTTCCAG AGGICIGIAC AAACACCAGC TITCATIGIG CITGGGAGIT
TCCATGCCTC TYCCITCTCT TCGCTTAGIG CACGITTCTG CTTTTTATCA GITTGACIGC CTGAGACTGA KTCCAACAAC
CCAAACTGAA CGCTCAGCTC CTCCKTTTCA AAGGAGGATG ACTINICINA ACAACTATIT AGGIGAATTA TIKCKACAGT
TTATTAAAGC AATGGCTCTA AACAAATTCC ACTGGGGGIG ACAAAGTACA ATACAAAAGG CGTACTCTGA GGGCTTGGGG
GT

SEO ID NO:42: (Length of Sequence = 278 Nucleotides)

AAACTITIGGC ATTITIATIC AGACACGIAT AAAAACAAAA CAAAAAACTI CAGIGATACA ACAGACGITI TCCCITAGIT
CCCCATCCAA GGGGACAGAG GIGIGCAGCI GAAGCIGGAY CITITITICIG TCCIACCIGG AAGCIGICIC ACIGCIGGAT
GAGAATGGCI TCIAAAAAGIG GATCITIGGGG ATCCITIGIGA ATTIGCCCIC GGATAAGGAG TGAAGWICAT TTACGGCACA
TGIGGATTAT GGITTACACA AAGATGICCA GITATITIT

<u>SEO ID NO:43:</u> (Length of Sequence = 225 Nucleotides)

AGATCAAAAG ATGAGAGAAG CTGAAACAGA ACCGCATGAG GGAAAGAGGA AAGTGGAATC TCTGTGGCCC ATCTTCAGGA TCCACCACCA GAAAACCCGT TACATCTTCG CCTCTTTTAC AAGCGGAAAG CCAGCAGCAG GATCTCTAGG AATATTAGTA TTAAAGAAGG CTATGCAGCA TAAACCTGAT TTCAAAATGG TAAAAGCAAG GTTATGTGTA CTTGT

SEO ID NO:45: (Length of Sequence = 305 Nucleotides)

GGATTGCCAG GAGCTGTTCC AGGTTGGGGA GAGGCAGAGT GGACTATTTG AAATCCAGCC TCAGGGGTCT CCGCCATTTT
TGGTGAACTG CAAGATGACC TCAGATGGAG GCTGGACAGT AATTCAGAGG CGCCACGATG GCTCAGTGGA CTTCAACCGG
CCCTKGGTAG CCTACAAGGC GGTGGTTTTG GGGGATCCCC ACGGCGAGTT CTGGCTTGGG TCTTGGAGAA AGGKGCATAG
CATCACGGGG GGACCGGAAC AGCCGMCTGG CCGTGCAAMC TGCGGGGGACT GGGATGGGCA AACGC

SEQ ID NO:46: (Length of Sequence = 264 Nucleotides)

ATGAAATAGC ATATCINNGC CTAATTAAAA GATTCCATTA CATTTACTTT TATCATTAT ACTGCCAAGG ATCAGTCACA
AAAAATTCAA ATTATACATA TTATTCATGC TTTAATTTCA TAAATAAGTA AATTAAAGCA AGCCAATATG TCTCTCTCA
TAACATAGGG AAAAATTACT GTTTAGCATA ACAGNGTAAT AGGCAAAGTC TAGCCATACA GCAGCAGTTC ACGGTGTTGT
CAAGTTGGKA CAGGTTCCAT CGAT

SE) ID NO:47: (Length of Sequence = 375 Mudleotides)

GATCTCTTCC AGCGTCAATG TACTGGGACA GCAAACACTC ACATTGAAG TTCCTTCTGG CCACCGGCTT CCCAGTACAT TGACGCTGGA AGAGATCATC TCAAATGGTT CTCCAGTGTC AGGCTGGAGA TCTCCAGAAA TGGAGTCTAC TCCTGGGGTG GCTTGTATGG GAGCC

SEO ID NO:48: (Length of Sequence = 270 Nucleotides)

GICIGICAGA GCNACOGGGC AGCICAMRCC CACAGCGGCI CCICATCCIC TGIGGIGGCA TCCTCATTCC ACICICATCI GCCACCIKCT CAGGCGGGCC TCIAGCITIC TCATGIACTC TAGCAATICC TGITTCICCT GCIGIAACTG CICCITITCC TTCIGGAGCA CACGCAGGC TGACCGCAGC TGIGICAGCT TCCGCTTACT TIMIGACAAC TGIACCAGGC TAGAATCCIT TCIGCCTGGG TCAGCTTCAG TCTTTGAACA

SEO ID NO:49: (Length of Sequence = 359 Nucleotides)

CCCTGAAGAG TGGGTGGGAC AACCAGATGG GTGTAACCCC TTGTGGGGGGA AAAGGAGTGA GTTTACTTGG TAAAATAATA
ATGGTAATGT CAGCAGCGTG GCTGGGGGAC TCAGTATGGT CCCGGGAAAA GAGTTGGGGC AGTGAACTTC CCAGGCCGAC
TGGCCTTGGG CTGGCAGCAG GGAGGCTGCA GGGCGCCTAC CTMCTCTGCC ACGTCCCTGC CTAGGAAACC TATCCCAGGA
CACCCTGCTT TGGCCTGGAT AGCAGCCTAG GGATGAGCAT TTCTTTGAAA GCAATTAGGT TATTCACCTG GTATTAAAAC
TATTTACTGT TAAAAAAATCT GTGACTTCAT GGARGTGGG

SEO ID NO:50: (Length of Sequence = 271 Nucleotides)

CCAGGAAGGA CAGGAAGTGT CCTCTAATAC GCATAAGATC CAGTACAGGA GAGATGGGAA GMGAGKCTCC AGGATGAAGG GGAAAARAGG CCGCATGCCA GTCACCTGGC ATCINCCAGA GAGGGYCAGY CINCCCACTG AGACTGGGGC ACGAGTCCCG TCATCACCAT GCCCTCTGAC TGTCGAACTG TCTTTTTACC TGACAAATAC TACACAGGTA TCGMTCGTGG CCATACTCTG CTATCTAAAC CCAGGAACTG ATTAGATTGT T

SEO ID NO:51: (Length of Sequence = 226 Nucleotides)

CTCCAAGCAG TAAAGACTTG CAAAGCATTG CATTTTGATT AAACCTTGCT GGGCTGAAGG GCAGGCAGAG CTGTGGTGGA CACTGGCAGG ACGCAGCACC CCCCGACTGG CCCTTGGCAG GCTGCACCGG GCGCATGCGG GCGCAGG GCGCTGCTTT AGGAAGCAGG TGGGAGTCTK NCACGTGCAG KCGGTCCAGG AGKGYACCAK GCCTGGCAGG GCACTG

SEO ID NO:52: (Length of Sequence = 408 Nucleotides)

GGTGGGGCAA GGTGGGGGTG AAGTGCACTC CTGCTGCATG AGTGGCAGGG CAGGGTGCAC ACACACACGT GGGTMCTGGC
TGGGTGAGGC AAGCAAAACC TGCCTGCACA TGGCAAAGGG ATGTGGGAAG TATCCATGGG CNCCAGGGGA AGCTGCAGTT
TGGGGAGGGA ATGGGTGGCA CTGCTGCGTG TCTGTGGGGG CCACCCCACT GGGGGTCTCC AAGTGGTCAA GTTCCGTCTG
CCAGGTTAGA AGCTATGATG GGGCCTTCTA GGACACTNGA GGCTGACCTG AAAGCAAGGT ACTTTTCACA CTGGGACCCT
GCAAGAGGCC AACAAGATTA AGGGATGCTT CAGGTCAGAC TTGGCCCTCT TCTTATGGGG CAAGACCTTC CCCGCAGAGT
TCAGATCT

SEO ID NO:53: (Length of Sequence = 314 Nucleotides)

TTCTGTGCAG GAGGACCACA TGGCAGTCCA GCAGACTGCA CATTITIANA AACTAGGTCT TCCCAGGTAG TTTGAGGAGC ACCAGGGCAC ACTCAGGGAA GGGACATGTC AGTGTCTGAG AGCTCACGGG AGGAAGGTGT AGTGACAACA TGGACCATGG TGGAGTGACT TTAGACGGCT CTTGGGTNAG GAGAATCATC ATGTAACAAA GCATTAAATC ATTTGGAGAA ATTCAGAAAA NTCGTAGATG TACATTCTAG CCCACTTACC AGGCCTACTA AACGTCAATC AGATATATTT CAATTTGAAT TCGG

.

SEO ID NO:54: (Length of Sequence = 310 Nucleotides)

AAGCCACCGC ACCTGGCCCA TTACATTTAT AATGITATAA GGGGGTTGAG GGGTCGTCCA CTGGAGCAGT GGTTCTCAAA CTCGTGTATG CATAGGAATT ACCTGAAGGG CTTGTTAAAA CACAAACTGC AGGGCCCACC CCCAGAGTTT CTGGTTGGGG AGGTGTGGGC TGGGCTTGAG GATGTGAATC TCTCACAAGC TCCCAGGTGA GGCTGCTGGT CTGTGGACCC ACTTCAAAGA CCCAGTGAAT CAGAAGAGTC AGTGAGACTG GACAAATGAA CGCAAGACAG TCTTCAAAGG AGACCAGAGG

SEO ID NO:55: (Length of Sequence = 252 Nucleotides)

TTTTTTTTT TYCCGGGGAR GICAAACATA CTTTTTCAAC ATAGGATKTC TGACAGGAGG CCCTTGGMCA GGGTTCCCTG
ACCTCTGYTT CAAACCCCAC TGGAAACAGA GCAAAGTCAT CAMGAAAACC CAGGACACCA GGGCAGGGGG GCTGCACAAG
GTCGGGTAGG TCACAGTGGG CCAGCACACA GTGGCCCCGC CCAGGTCCAG CCCAGCCTGG GGGAGGGTGT GAGGGTTCCA
KGCAAGCTCA TT

SEO ID NO:56: (Length of Sequence = 188 Nucleotides)

GTCAAGTCTA CCATCATTCT AGAAGGAAAA GGCATGGTGG GAATTCAGCA CCTGAACTTG TATTTACACC AGCCTCGGCA TCTGGCAAGG RAATAGCGAT TGTTCATAGT GATGCAGAGA GAGAACAGGA GGAKGAAGAA CAAATACACA CAAACAACTG ATCTAGGGAG ACTCCAARGA TCCAACAG

SEQ ID NO:57: (Length of Sequence = 304 Nucleotides)

AATCAGCCTG CAAGCAAAAG ATAGGAATAT TCACCTACAG TGGGCACCTC CTTGAAGAAG CTGATAGCTT TTACACAGTA
TTAGATTGAA ATAATGGACA GAAACACATT CTTGTCAAGA AAGGGGGAGA GAAGTCTGTT TGCAAGTTTC AAAGCAAAAA
GCAAAAGTGA AATGATTTGA GGATTTCTGT TCTAATTGGA GATGATTCTC TGGTTGTTAG AAATGGCAAA TATTGATGAT
TGTGGGCTAT TGATTGGTGC AGGATACTTG GTATACGAGT AAATACTTGA GACTCGTGTC ACTT

SEQ ID NO:58: (Length of Sequence = 261 Nucleotides)

CCAGAAGCIT CITGCCTTCT CIGIGCICTC AGIGGTTCCC TICCCTGAAG TGCCTCCCTT CICATTAATT ATAGCCTGTG
TCTGAACATT GIGAGCTATA AGAACCCTCA TATTAATGGT TAAGGGACTG TTGGAAATGA TGTGATTTTA TTAAAAATGG
GGTCTTTGTG GAGGAGTCAG GAATGGTCAA AATGAGCTTC AGGTATGGGG CITGCTCTRT GCTCCTGATA CCAAGGGTCT
GGCAAGCACA AAGGAAGGTG G

SEO ID NO:59: (Length of Sequence = 470 Nucleotides)

AATACGIATT CIGAAGCCAC TATATCIGCA TATGIATCCC AGATITGAAC AATIAAGTAA AAAGATGGIG AATGATGAAA
GCCAGITITC TGICIGIAGA AGTGAGAGGI GACAGATAAC CAAAGGAAGA AGGCTAGAAT GGATAGAGGA CAGTGCTTAA
GTGTAGITCC TGITGCCTTT AGTCTTATAG ACTTCATTIC CAAAGTTTCT TAGCACCCCC CITCCCCCTT TGGTGAGGIT
GTTTCACATA TTTTCTAGAC AATTAGATTC TTTTGTCAAA GTCTGTGTC CATCCGGAGA GCCTCTGATC TCTTAAATGA
TTTTTTAAAT TTACATACAT TAAGGITCAC TCTGCTGTAA AGGTCTGTGG GTTTTAATCC TGTCTCACAG TTTTTGCATA
TGTTGGCCTT CTGCCTGGGA ATACTCTCCC AGATATTCCC CATGACTGGC CCCTTATCTT CAATCAGATC

SEO ID NO:60: (Length of Sequence = 466 Nucleotides)

GIGITICAAG GGAAGGCAAC IMCAAGITIG IGCAGCIGAA ITITCIGIAAA GITAAGACAG ACTCAMCITC ICATICAATC IGGGGCAGIG GATAACCITT CIGAATAGAC CCACITGIIC ACGGACAGGG ATAGAGGIIT GCCITICIIC ITITCCITGAA ITITGGAGIGA GCACITAGGGA GGGGAAGIGC ATGGGIGACA IGAAGAGGI GAAGATGIAG ITAAAAGCATC ATCCAGGIAC ACAITAACGC IGCCCAGAAAGC INTITACIGAG CACAAAAGCCI

AGICCICAAG GECIGATICC ACCITCCCIG TCCAGGGACT TICTCAGCAA ACTITGITCA TGAGCAGIIG TICGCITIGA TGGICTIAGC CAGITTITGG TGCAGGGGIG TICCICTGGI ACTAGGGCIA GGGCAGCIGI TIAAAG

SEO ID NO:61: (Length of Sequence = 491 Nucleotides)

GACACCCCTC CIGCCATGAA GAATGCCACT AGCTCTAAGC AGCTCCCACT GGAACCAGAG AGCCCCTCAG GGCAGGTCGG
GCCTAGGCCA GCCCCCCCC AGGAAGAGTC CCCTTCCTCT GAAGCAAAGA GCAGAGGACC CACCCCACCA GCCATGGGCC
CACGGGGATGC CAGACCTCCT CGAAGGAGCA GCCAGCCATC TCCAACAGCA GTGCCAGCCT CCGACAGCCC TCCCACCAAG
CAAGAGGTGA AGAAGGCAGG AGAGGAGACA AAGCTGGCAA AGGACCGGCG AGAAGAAGCCT CCCAAGTIACC TGGCGGCCAA
GGAAGGCAGT GTGGCTGGGA AGGAGGAGAA AGGCCAAGGT GCTGCGGGAG GAAGCAAGCT CCATGGAGCC CCGCTGCCGG
TTTTAGGGAG CAAACGTCTT AAAGCCGAGC AACGCCGTTC AAGCCTTGGA GGAACGGCTA GCCGAAGAAG TTTTGTGGAAA
ACAAGGGGCC T

SEO ID NO:62: (Length of Sequence = 478 Nucleotides)

ATCATTGAGT ACGCAGAGCT CAAAACAGAC GTGTTCCAGA GCCTGAGGGA AGTGGGCAAT GCATCCTCTT CTGCCTCCTC
ATAGAGCAAG CTCTGTCTCA GGAGGAGGTC TGCGATTTGC TCCATGCCGA CCCTTCCAAA ACATCTTGCC TAGAGTCTAC
ATCAAAGAGG GGGACCCTCA GGAGGTCCGG ATGAAACGTC TGGAAGCCAA GTATGCCCCG CTCCACCTGG TCCCTCTGAT
CGAGGGGCTG GGGACCCTCA GCAAATCGCC ATTGCTCGCG AGGGTGACCT CCTGACCAAG GAGCGGCTGT CTGTGGCTGT
CCATGTTCGA GGTCATCCTG ACCCGATTCG GAGCCTACCTT CAGGACCCAT CTGGCGGGGC CACCGCCACC AATGCGTATG
ACGTCGATGA GTTTTTGAGT TCACTGCTGT GAGCGCATGA GTCGTGTACT GAATCCTGTG GACAACGGTT AAGTTACA

SEO ID NO:63: (Length of Sequence = 183 Nucleotides)

CCTGGAAAGT GGGGTGGGC CAGGGGGCCA GGCCCAGCAT GCACCCCCAT TITTITIGGG GCTGATCCCT GCCCCAGCTC TGCTGATACC CGGGGCCACA GCGTCAGGCC GTTGGGGGTG GAGKTAGAGG TGGGAGAGCA GGGGAGAGAG CCTKAGGAGC CACAATTGGG CAGACAGAAG CGG

SEQ ID NO:64: (Length of Sequence = 316 Nucleotides)

GGATATIGCA CCITACAGAC TIAGGGAGCC TITACCAGAG ACGCCIAAAA CGCCCCAGGI TCAGCCATIG TGCIGAATAG
AGIGGAATAT AGAACCAGGG ACAGAGIATI TCATITAACG TIGATATATA CITGCIAAGG AAACACTAAC AATACIGIAA
CITTGITAAA GGACATAGIA TIGAAATGGG AAATAGAGGI CAGGCICACA TCATCITAGI TIAATGCIGG GCAACITTIT
CIGATTICIG TAGTICCCIG GAAAATGIGI CCITCGIACC CATAAAGTGG TACAAATGCA TITGIAACCA TITTIG

SEO ID NO:66: (Length of Sequence = 411 Nucleotides)

ATCIGGICIA GAGAGGOGAC TOCAAGCTOT CITGCIGGCT COCAGCIGIG GGAATCCITT AGGCIIGITC TOAACCIACA
CGITAAAAAT GCITCITGGT GIGITIGGGG AGGGGGAGAG GGAAACIGAG CICICICITG ACCTOCTOCA ACACCCITGA
CTIGCITACC CAGCCATITI CAGIAGCIAC ACGGGIGGIC ACAGAACACT GGGCGGCACT CGGCACACAA CACAGAACCG
GGGCAGICCA TGCAGGIGCG GGAACACATG TCGGACCCAG GGAGCAAGGA ACACGCCACC CCGAGGAACA TGCAAACCGA
GGAAGGATTC CCTTCAGATT CCAAGGATGC CACAACCCCG ACGGGCGGCT TAGGGAGGCA CCGATTATCT AAGGAAAAAG
GCCACTGTIT G

SEQ ID NO:67: (Length of Sequence = 413 Nucleotides)

CIGCICCITA TGITTITATI TCCAAAGITI AGAATITCII TGCITCATAG TATIATITIA TITTACIAAA TIACAGAGIA AGAAAAGCTI TICATITTAT CIGATITIAT ICITAGAACA AAAATATTAC GATCITCIAT ATATITGTAT TITTIGCCAAA AASTGTAGGC AATTITACAT CATCITTITI CCCAATCAGT TIGIGATCCA ACTATAAAAA GGAGACATAG AATACIGAAT AAATGAAACA GAAACTCCAA GGCCAAGAAG TGTCCATCIT GAAAGAGTGT TAGTGGCAAG ATATGTGACT GCAGACTAGA TGTAGACAAA CCTGAGAAAA ACCAAGCATG GGGGAAAGGA TYCCTATTIT AATAAATGGT GCTGGGGAAA ACTGGCTAGC CATATGTACT TTA

SEO ID NO:68: (Length of Sequence = 372 Nucleotides)

GCACGGTTAA AAGACCAACG TGTGTGGNTC AAATATAAAG GCCACACCTT TCAGACCGAA CCTACTCAAA GATCCTTTAC

TTTGCAATAA TTTGAACTGG AGAACCAAAG ACGGGAGACG AATGAAAGCA AAGATCCTCA AAGAACCAAA GGAAAGACCT

GAAGGAATCC ACCTGCATAG GCCACGCGTT CCACTCTGGG TCAAATGCTT CCACGATGCA GAAACCTTTT TTTAAAAAAAG

TGCAAGTCTA ATTACCTACC AAGGGTAATA AAAAGCACAG CACAGGAATG ATTACAGCTG ATGGTCAAAA AACAAACCAA

AACCATTAAA AAAACAATCA GGCAGAAAAC AGGAGTTAAA TGTTTACATA TG

SEO ID NO:69: (Length of Sequence = 389 Nucleotides)

TCTAGAACCT GGACCCACCC AGCGCGTCCT TTCTTATCCC CGAGTGGATG GATGGATGGA TGGATGGTAG GGATGTTAAT

AATTTTAGTG GAACAAGCC TGTGAAATGA TTGTACATAG TGTTAATTTA TTGTAACGAA TGGCTAGTTT TTATTCTCGT

CAAGGCACAA AACCAGTTCA TGCTTAACCN TTTTTTCCTT TCCTTTCTTT GCTTTTCTTT CTCTCCTCTC ATACTTTCTC

TTCTCTCTCTT TTTAATTTTC TTGTGAGATA ATATTCTAAG AGGCTCTAGA AACATGAAAT ACTCAGTAGT GGATGGGTTT

CCCACTTCTC CTCAATCCGT TGCATGAAAT AATTACTATG GTGCCCTAAT GCACACAAAT AGCTAAGGG

SEO ID NO:71: (Length of Sequence = 329 Nucleotides)

GAAAAAATGG GAGGGCAGCC ATGITATTAAT TGITACATCCA AGGAAACTGT GCCCCAGGGG TCITGIGIGI ATITCTGAGA
AGAGGGGTGA GAAAAGGCAC TGIGICAACA TTIGCITCTG CCIGAACGTG CACCTCCCAG TGCTCCTCCA TCAATTAGGA
GAACTGICTT GAAGAATGCT GCCTCAGCTT CTGAAGAGAA GACCCCAGGA CATGCATTAA TGAGAGGAGG GGAGTCACAG
CTGCAGAAGA ATAAAGCTCT CTGAGGGAGC CTGGCAGCCC CCAGTGGAGG CCTGGAGCTT GTTGACCANN GCAGCAGGAG
ACCCCTGCT

SEO ID NO:72: (Length of Sequence = 418 Nucleotides)

CTGAGTTGCC TGAGGTCATT CACATGCTTC AGCACCAGTT CCCATCTGTT CAGGCAAATG CAGCGGCCTA CCTGCAGCAC

CTGTGCTTTG GTGACAACAA AGTGAAGATG GAGGTGTGTA GGTTAGGGGG AATCAAGCAT CTGGTTGACC TTCTGGACCA

CAGAGTTTTG GAAGTTCAGA AGAATGCTTG TGGTGCCCTT CGAAACCTCG TTTTTGGCAA GTCTACAGAT GAAAATAAAA

TAGCAATGAA GAATGTTGGT GGGGATACCT GCCTTGTTGC GGCTGTTGAG AAAAATCTAT TTGATGCAGA AGTAAGGGAG

CTTGTTACAG GAGTCTTTGG AATTATCCCT CATGTGATGC CTGTAAAAAAT GACATTCATT CGAGATGCTC TCTCAACCTT

AACAAACACT GTGATTGT

SEO ID NO:73: (Length of Sequence = 336 Nucleotides)

CTGAATTITT ATATECTICA CTTAGGCTTT CATTIGAGTA GACTCTAAAA ATTCTGCCTT GCTTAAGINC TAACACTGCC

TCTCAGATTT CAGTTTTGGA CATTGCACAA CTAAGACCTT TTAAACGCAT TINCTTGCTA ACTCGGAAGA CACATAGTCT

GCAGCAAGAC ATTCCTATAT TGAAGAAATG AGAGAAAATT TTATGCTGCA TCAGGTGGAG AGCAAGGCTC AACGGTGGTT

GCATTAGTTC CCTCGGAAGT ATTGAAAAAN CTTTGAAATG GGAAGGAAAA TTTTTTGCAC CTAATGTTCC TGAGGTACCC

AGAATGTCTG GGGGTT

SEO ID NO:74: (Length of Sequence = 402 Nucleotides)

GIGCTCAGIA AATACAAATT GGATGGACIA GAGAGATAGC CCCGAGGACA CTGCCAAATA AATAACAAAT TGIGCAAGCA GCAGGCCGCT GTAATTAGAC CAAGGAGGAC AGTCAGTTAT TAATATCAGA CACGTGGCAG GGTTAACAGC CACTGAGGGT GGGTACAATG AAGAGAGTCA CTTTCTGCAC CCTCAGGGAC TTCCCTTGTG ATGGCCTTCT AAAGAGGGCT GAACAGCACC AAGTGCCCTC GCTGCCTCTG GTTCCTGCTG CCCTCCGCGT GCCTTGGGTG CCCCACAACT AGGGCCCTGG GTCCCTCCCA TGTCCCCCTC CCTCCTACAA CCCCTCAGCC CCTTATCTGG CCAGCCATTA TGATGCCTAT CAGTATGAGG CCAGATGAGA GT

SEO ID NO:75: (Length of Sequence = 454 Nucleotides)

GEACCCCGGG CCCGCGATGT GGCCCAGTAC CIGCTCTCAG ACAGCCTCTT CGIGTGGGT CIAGIAAATA CCGCTTGCTG
TGTTTTGATG TIGGIGGCIA AGCTCATCCA GIGTATTGIG TTTGGCCCTC TICGAGTGAG TGAGAGACAG CATCTCAAAG
ACANATTTTG GAATTTTATT TICTACAAGT TCATTTTCAT CITTGGTGTG CIGAATGICC AGACAGTGGA AGAGGTGGTC
ATGTGGTGCC TCTGGTTTGC CGGACTTGTC TTTCTGCACC TGATGGTTCA GCTCTGCAAG GNTCGATTTG AATATCTTTC
CTTCTCCANCC ACCACGGCGA TGAGCAGCCA CGGGTCGAGT CCTGTCCCTG TTTGGTTGCC ATGCTGCTTT TCCTGCTGTG
GACTTGCGGC CGTTTGCTCA TTACCGGGTA CACCACGGAA TGCACACCTG GCTT

SEO ID NO:76: (Length of Sequence = 313 Nucleotides)

GCITIGATAG CIAGTIGICI AAAAGIGCIG NITATIAAAT AATOCACCIN TITOCOCACI TAAAACATOC CICTIACCAT
ATACIAAATI OONGIAGOOC TGGGICIGIT TCIGGACICI COCGICIGIC TGACCOCCIC CAGGICACAC TGAGIGAGGI
AATGGIGGOG TGAGAATOCI CIGGGAATCI GGCAGGNICA COCCNGAGCA GICCACCCCN CAACICATTA NCATCGITCA
GAGIGGNCIG AGIGNICICA CACATICACI CIGCCAAATG CACITIAGGA ACTGICAAAT TCCAAAGIIT CAA

SEQ ID NO:77: (Length of Sequence = 446 Nucleotides)

CTCAGCOGIA GCCCTAAGIC GITTITCCAA TITAGGAAGC TCACAACGCA GATCIGCATT GICACGIACC AGCIGITIGI GAACCITIGI AAGCIGITCC AGGITGITCI CAAGAAAGGA AATCITCIGC TITIGGGAGT GAATCCCCCC ACTGICITCG GGCTCCATTT CIGCACITIT CITGACICGA GICGIGACGT CITGAACGAA CAGCITGCGA AGGITGIGGC SGGICTGGAG TTCCCCGGGCA ACTGICTCCT CCAGACCCIT GAGGICCTGC TIGIGACTGC TCAATGICGC TCGIACAGAA ATGICAGCIC CIGCAGCITT GGIGCICTIC TCGICGITCT AGCICTITC AGCITTCTCG TAGICAAGCC TGAAGGCTIC TCTAAGCICT AACTGGAGCT TCTGATTTAA GGICTTITGA GCICATCAAA TGGICT

SEQ ID NO:78: (Length of Sequence = 296 Nucleotides)

ACCOSTISCE GEAATGEAGA GAATGIGCCT GAGACAGAGE GCCTGGCTGG GGAGGAGGAG GCCCTGGGRG COGAGCTCTG
TEAGGAGACC CCTGTGAATG ACAACTCATC CATCGTGGTG CGCATCGCGC CCGAGGAGGAG GCAGAAATAC GAGGAGGAGA
TCCGCCCGTCT CTATAAGCAG CTTNACGACA AGGATGATGA AATCAACCCAA CAAAGCCCAAC TCATAGAGNA GCTCAAGCCAG
CAAATNCTGG ACCAGGAAGA GCTGCTGGTG TNCACCCGAG GAGACAACGA GAAGGT

SEO ID NO:79: (Length of Sequence = 285 Nucleotides)

CCTITICCTGC CTGGGAAGTG ATGACTCGCA GGTCGGGCTT GCGGCTGGGG GCTCCAAGCT GGGTGCTGTG GGTAGGTGGG GGCGGAGACT TGGCCAGGGAT GACCTTGTTT AGGCTGTTGC CATTGGCCAC AGGGAGGAGG CCAGGGGAAG CCCGAGCACT GACGTAGCCA TTCCCAACAG GGCTGGGGCA GGCTCCGTTA GCACTGTTCA GGTCACCNCC CAGCATGGCC CCCCCACTACGCTG GGGCAGGCCA GGAGACACAC TGTTCCTCTG TAGTG

....

SEQ ID NO:80: (Length of Sequence = 402 Nucleotides)

ATGATTICTT GCCTGINATA ACCTATGCAC TCACAAAGAT GAACTCTCTG AGAGGGATGA GCAAGAGCTT CAGGAAATCC GAAAGTATTT CTCCTTTCCT GTATTCTTT TCAAAGTGCC GAAACTGGGC TCGGAGATAA TAGACTCCTC AACCAGGAGA ATGGAGAGCG AAAGATCACC GCTTTATCGC CAGCTAATTG ACCTGGGCTA TCTGAGCAGC AGTCACTGGA ACTGTGGGGC TCCTGGCCAG GGATACTAAA GCTCAGAGCA TGTTGGTGGA ACAGAGTGAA AAGCTGAGAC ACTTGAGCAC ATTTTCTCAC CAGGTGTTAC AGACTCGCCT GGINGATGCA GCCAAGGCCC TGAAACCTGG TGCACTGCCA CTGCCTTGAC ATCTTTTATT AA

SEO ID NO:81: (Length of Sequence = 246 Nucleotides)

CATTITIAAT AGAGACGGG TITAACCATG TITGCCCAGGC TGGTCTTGAA CTCTTGATCT CAGGTAATCC ACCCACTATG
GCCTCCCAAA GTGCTGGGGT TACAGGTTTG AGCCTCTGIN CCCGGCCCGG CCAAAGACTG CCTATTCTAA ACGTTGCTGA
GGACGTGGAN CAATCACAGC TCTCCTNTCT TTCCAGTGGG AGTTTAACAT GGCACAACCG CCTGAAAACC GTTTGGNGAT
TTCTGT

SEO ID NO:82: (Length of Sequence =394 Nucleotides)

GGGAACCCTC AGCAAAATAT AATGGTACCG CTATTATCAG CCTTGTTCGA GGCCCAGGGA TITTGGGGGA GGTCACAGTG
TTCTGGAGGA TATTCCCTCC TTCCGTGGGG GAATTTGCTG AAACATCAGG NAAACTGACA ATGCGAGACG AACAGTCTGC
AGTCATTGTA GTAATACAGG CTTTGAACGA TGACATTCCC GAGGAAAAAA GCTTCTATGA GTTTCAGCTC ACTGCAGTCA
GTNAGGGAGG AGTTCTGAGT GAATCCAGCA GCACTNCCAA CATCACGGTG GTGGCCAGCG ACTCTCCCTA TGGCCGATTT
GCCTTTTNAC ATGAGGCAAC TTCGAGTGTC AGAAGCACAG AGGGNTAACA TCACAATCAT CCGTTCCAGT GGAG

SEQ ID NO:83: (Length of Sequence = 308 Nucleotides)

SEO ID NO:84: (Length of Sequence = 313 Nucleotides)

CTTTAACTTA ATGGCAATTA AAACTCACTG GCAAAAAAA TCACTAGAGA TGTCAGTCCA TTATCTTACC AAATAGTGTA
TTTTTACCAT CTTTTACCTA CACCCTTGAG TAAGGTGGAA TAGGTTAAAG TTACTGGCAT AATAACACTT CATTGAATTC
ATGATAGTAT TTAACATGTT AAAACTGTTT AGTTGAAAAG TTCACATGCA ATTTATAATT TAAAAATATG CTACATATAT
TTCATAAAAAW TACAATAGGT CATACTARAC TTTGACTAAA ATTAAGAATG TKTTTCTKTC ATAATAATGC AGG

SEO ID NO:85: (Length of Sequence = 303 Nucleotides)

TECTCOSTIT ATTECTCTAT TCAATGACCA CGAGCGAATT ATAAAAAGAC ACCAAATGTC TCTGTCTGCC GTGGGATAAA
TATTTAAAGT CAGCAATAAA GTCACGTGGC TCCAAGRTAA TACATGTTGC CAAAGAGTCA TGCATGCCCT CCTGATGGGC
TCTCAACACA CGTATGGWCA TGGGAACACA CGCAGAGCAA CACGCAGTAT GAACITSTGG GAAGGCTTTA CCACAGTGAC
ACAGTAAAAT GTCTCACGTA GATCTGRGCT GAGTCCCCAC CCAAACCTTG AGCTCCCCTT CCA

SEO ID NO:86: (Length of Sequence = 380 Nucleotides)

SEO ID NO:87: (Length of Sequence = 280 Nucleotides)

GCCTTTGCTG CTTATTCGCA TCGATGGTGA AAGAGATGTC AGGAGCACTT CTCTGCTGAG GTGGCTGAGA CGAAGAGGAC
TCTGCTGCCCA GCCTTGCCGC ATACCTGGCA ATTAGCCTGT GTTCTTCATC AAGCCGGTTT GAACTCTCAA GCATGCTCCT
GGTAATAAAA GGACTTCCTG AGGAGGGAAC AGAGTGNEAG AACAGGGTGT CGTTCATGCT GGTTACAGGT CTGGGAGGCA
CGATGTGAGC CAAGTTGAGT GGCTTCTCAG GCTGATCTGG

SEO ID NO:88: (Length of Sequence = 446 Nucleotides)

CCTGCGTCTC TTACACCCYC TCCCACCCGA GGCTCCCCCGG AGATAGCAGA GAATTCGAAG AGGTCGCCGG GGACTGGAAA
GAAGTCCCNG NAGGCCGCCT TCGCAGTCTA CACCCCAGCC TGCTTCCCAG CCTACAYCCA GACCCAGCTC AGACCTTCGT
GACCACCCCA TCCCTTTCTC CGGCTGGCTG GGTCGGGGGC ATCCCTCTCT GTCGCTGGCT TCCAGAGGCA GGACAGGCCT
CCTGGTAAGC CCGCAAAGTT GCTGACCTCC TGACTTCGTC TGCCTTTTAT TAATATCTGT ATTGCTGATA ACCGTGCTCT
TGACTATGTG TCCCAGGTCA TGTCCCAGGT CATGGAGAAG CCCGTGCCAC AGTGACCCTT CCCATACTTC TGGGGGGGCT
GCTCTCCATC TGGATCGTAG GAGGATATAG GTGTGTTCTG GACCAT

SEO ID NO:89: (Length of Sequence = 384 Nucleotides)

GROCCITCIG GGGACICIRI TICCCCATIT ATRECTECIG TGTCCCINAC CAGTICCTIG CAGGATICCC TCCTITTAAA
ATGCCCTIAA ATCTAGCTIT GCCTIGGAGA CCCCAGTGGG TGCTGCTCCT GCCGTTTTCT TCCTGCCAAG CCTGAATCAA
TGTTTCATCT CCAACCCTCT GCCAGTTTGG CCCCTCAAAG CTTGGTGGCT CAAGACTGTW AGCCTGGCAG AGCCGCGAGG
TGAAGGGAGA AGCTCTTGGA GCAGGCAGGA TGCCACCGCT GCTTCAGCTT GCCTCCTCGC CCAGCTACCC TTTGGCCCCCA
TTGGGCCCCTC GIMIGCCTCT CCAGGATTGT ATGTTTCAAG NCTTGTCCTG TGTTCCTTTG TCTG

SEQ ID NO:90: (Length of Sequence = 344 Nucleotides)

TCAAGCTGGA AAGGGCTACT ACCTCATGCT GGAAAGGGCT ACTACCTCAA GCTGGAAAGG GCTACTACCT CAAGCTGGAA
AGGGCTACTA CCTCAAGCTG GAAAGGGCTA CTACCTCAAG CTGGAAAGAG CTACTACCTC AAGCTGGAAA GGGCTACTAC
CTCATGCTGG AAAGGGCTAC TACCTCAAGC TGGAAAGAGC TACTACCTCA AGCTGGAAAG GGCTACTACC TCAAGCTGGA
AAAGGGCTACT ACCTCAAGCT GGAAAGAGCT ACTACCTCCA AGCTGGAAAG GGCTACTACC TCATGCTGGG AAAGGGCTAC
TACCTCAAGC TGGACAGGGC TACT

SEO ID NO:91: (Length of Sequence = 364 Nucleotides)

GOCCCAGGGT GAGGGCTATG AGGGGTCAGG GGTCAGGTTC CCCAGGACCC TAGTCCTTGT CCCCTTCCCT GGTGCTAAAT
AAAAGTGAAT AAATACTAAA TAAATACAAC TGGGGCCCAG GCCCTCCCTG CCTTCCCCCT CCCTCCTGTG ACCCGCAGCA
GAGGGGGCAG TTTAGATGGA GGGCTGTCTG TCAGCCCCTT CCCATCCACTA ACCCATCACT GCCTCCCAGG GCAGGAAACC
AGGGCAGGGC CAGCCTGCGC ATTAGGGCAG AGAGGAGGGG CAGGTCTCAC GCCCACAGCC CCTTCCCACT TGAGTCTTAG
CATGAGGCAG CAACAGAAGC TCTCTCTTCC TCCCAGCTAA GTCC

SEO ID NO:92: (Length of Sequence = 218 Nucleotides)

ATTTAATAGA AAATTAAAAT AATAAATAAT ATGAAAAAGA :.TGATAACGC TGAGCTGGGC AGGCCCAGGC CAGTCTAGTA
CAAAGTTAAG GAGGTAGGGA GGATGGTGGG GAGGAGGGGG CGGACTACCC TGCAGGACGC GGGAGGCTGC TCAGACTGTG
GTGATGTCAG GAAGGGCCGC ACACTTTGGC ATGGACGATG CACTAAAAAA AGAGAAAG

SEO ID NO:93: (Length of Sequence = 364 Nucleotides)

GCTTTCAAGG GAACAAAGAA TOGGCCTGGC AGTGCCCTGG AGAAGGAGGT GGAGAGCATG GGGGCCCATC TTAATGCCTA
CAGNACCCGG GAGCACACAG CTTACTACAT CAAGGCGCTG TCCAAGGATC TGCCGAAAGC TGTGGAGCTC CTGGGTGACA
TTGTGCAGAA CTGTAGTCTG GAAGACTCAC AGATTGAGAA GGAACGTGAT GTGATCCTGC GGGAGATGCA GGAGAATGAT
GCATCTATGC GAGATGTGT CTTTAACTAC CTGCATGCCA CAGCATTCCA GGGGCACACC TCTAGCCCAG GCTTTGGAGG
GGCCCAGTGA GAATGTCAGG AAGCTGTCTC GTGCAGACTT GACC

SEO ID NO:94: (Length of Sequence = 423 Nucleotides)

CITCATACIA GAACIGICIG CCATCITIAT TICTITGITI TCAGGAAAAT TGGAGAGAAA AGIATITCIT TITTAAAAAT
GATTATTATA CITTAAGITC TGGGATACAT GIGCAGAACG TGCACGITIG TTACATAAGI ATACACGIGC CATCGIGGIT
TGCIGCACCC ATCAACCCGI CATCTACATT AGGIATITCI CCIAATGCIA TCCCICCCCT AGCCCCCCAC CCTCCAACAG
GCICCAGIGI GIGATGITCC CCICCCTGIG TCCATGIGIT CTCATTGITC AACTCCCACT TATGAGIGAG GGACATGCAG
TGITIGATTT TCTGITCCIG TGITACTITG CTGAGAATGA TGGCTTCCAG ATTCATCCAT GICCTTGCAA AGGCATGAAC
TCATCCITIT TATGGCTGCA TAG

SEO ID NO:95: (Length of Sequence = 405 Nucleotides)

AACAGCCCCC GATCIGCATA GCCTGTGAAA GCCCACGGGG ACATCAGTAA CCTTCTGCAG CCACCATCCA ATGCCATTAC
TGINAAGIGA GACTIGGCCA CTGIAGCCTG GGCCTGCTGC AGGAGCTCTT CAGAAAGGCA CATGAGGACC ACGGTTTGCC
TCAGITTCTG GTAAAACACA AGGTCTGGAG TGCCCCTGCA AAGGGTATTG ATGGACTTCC TGCCAGTGAC AGAGCATGTC
TATTGCAAAC AATTCTCTCA GTTACGTTCA GCACTTAAGA ACGGCTAATG NCAATAGGAT CTTTAGCAAC TTTTTCACAT
CATAGAAGGT GCAATCGCTC ACTTGGGAAC ACTACTGAGA GTGACTTCTC TTTTAAAATT GAGTAGCAGA TGAAAAATTA
AAATT

SEO ID NO:96: (Length of Sequence = 173 Nucleotides)

GAAGACAATA CTGATGCCAG CTCTTTGTAA TTGTGAAATC TGTACCCAAA CCTCTGGATT AGAATCTCCA GTTGTCTACT GTAAATACTG GAATTACAGC AAAGGATATG GGGACTGGGC TGCTTTTCTG TATTGTACAA GCACTATTCT AGATATTAAA GAAATTTAAC CGC

SEO ID NO:97: (Length of Sequence = 337 Nucleotides)

ATGGCGCCCT ACAGCCTACT GGTGACTCGG CTGCAGAAAG CTCTGGGTGT GCGGCAGTAC CATGTGGCCT CAGTCCTGTG
CCAACGGGCC AAGGTGGCGA TGAGCCCANIT TGAGCCCAAC GAGTACATCC ATTATGACCT GCTAGAGAAG AACATTAACA
TTGTTCGCAA ACGACTGAAC CGGCCGCTGA CCCTCTCGGA GAAGNTTGTG TATGGACACC TGGATGACCC CGCCAGCCAG
GAAATTGAGC GAGGCAAGTC GTACCTGCGG CTGCOGNCGG ACCGTGTGGC CATGCAGGAT GCGACGGSCC AGATTGGCCA
TGCTCCAGTT CATCAAG

SEO ID NO:98: (Length of Sequence = 212 Nucleotides)

TGAAGCCCAA GNAGINIGIG AAGACAGAGA ATGACCACAT CAACCIGAAG GIGGCCGGGC AGGACGGCTC CGIGGIGCAG TICAAGATCA AGAGGCACAC GCCGCIGAGC AAGCIGATGA AGGCCTACTG AGAGAGGCAG GGCTIKTCAA KGAGGCAGAT CAGATTCAGK TICGACGGGC AGCCAATCAG TGAAACTGAC ACTCCAGCAC AG

SEO ID NO:99: (Length of Sequence =: 26 Nucleotides)

CCTTTTAATA ATAATTCTGC TGTCTGCTGT GTACTAGAAC CCATGCCTAC TGCTTGGGGT ATAATGTAGT AAATGTAGTA
AAAACAATAT CCGCCGGGGG CGGTGGCTCA CGCCTGTAAT TCCAGCACTT TGGGAGGCCA AGGAGGGCGG ATCACGAGGT
CAGGAGGCG AGACCATCCT GGCTAACATG GTGAAACCCC GTCTCTACTA AAAATACCAA AAATTAGCCA GGCGTGGTGA
TGGACGCCTG TAGTCCCAGC TACTC

SEO ID NO:100: (Length of Sequence = 333 Nucleotides)

AAAATGCTCA CAGTGGTCTT CTCTGGCCGG TGAGCCTACA GCTGATCTTG TCAGAGACAA ACGTTAGTTT TACTGAGTCA
CCCAGAGCCC TGTGCTGGTG CCTGAGGGTT TGTTCCATGG GACAGTCTCC ACAATTCCTC TGGGGAAGGG CCACAAATCC
CACAGTGTGT CCCCAAGAGGG CTGGAGTAGG CGGAGTCCCC AGCAGCTGTG GCATGACCAG CCATCTCTCT CAAAACAATT
GTTAACAAGC CTTCTGCAAG TTAAGGTTCC ACATGGTAGC CGTGGTACAG AGGCATTTCT CTAGGGTGGG AGAGGCTTGT
GCTCTACACC AGG

SEQ ID NO:101: (Length of Sequence = 156 Nucleotides)

CTCTGACTTT CCTGTGGNTT TAGAGCCAAG CTCAAGGTAG TAGGCCGTAG GGNCTTATTT TATTTTCAAA CCCCCATCCT CAGAGCGCAG ATACATGCAG AGGCTTCTGC CAGGCTACCA CGGGGCCTTA GTGGGAACAG GTTGAGACCA GCACTT

SEO ID NO:102: (Length of Sequence = 331 Nucleotides)

CGAAAAGGG NNVIATGGCC ATCITITATC AGAAAAAGIG ACAAAACGG AATTIAAAAA ATGAATTITC NNTCTGACTT
TATTINNAAA TACACITICI TITTINNAAA ACCAATACAC TITCITTGAG GATGACAGIA TTAGGAAATC CAATINNACA
AAAAATACTA CATCIAGICI GGGGIAGATA TATTIATTIT TGGIAACATA CATTAAGIGG CACTAATTAC ACAGTAACTA
TAAGGIAACT AACATGAAAC CACAGAACTG TAACICTGCC ACAGCTGCAT GAACITGGGC TITTCTGGIT GAGCCCATTT
TCAAAAAAACT G

SEO ID NO:103: (Length of Sequence = 316 Nucleotides)

AGCCACTGCG CCCCACCCCA TITOGGIGIN ANCICAGCTC ACTICAACCT ACCCCTCCCA AGITCAAGIG ATTCTCCIAC CTCAGCCTCT TEAGTAGCTG GGATTACAGG GGICTGCCAC CACGCTGGGT GATTTTCCTA TITTTAGTTG ACACTGCATT TCACCAGGTT GGCCAGGCTG GTGTTGAACT CCTGACCTCA GCTGATCCAC CCGTCTCGGG GTCCCAAAGT GTTGGGATTA CAGGTGTGAG CCACCACACC AGGCCCATAT TTTCTTTTAG ACATGCAGGC AATGTTGGTG GGTTTGTCTG TTAAGA

SEQ ID NO:104: (Length of Sequence = 308 Nucleotides)

GITTITICCIG CATCIATICA GATAATCAIG IGGITTITGI ATTIGGCICI GITTATAIGC IGGATIACAT ITATIGATIT GCGITATAITG AACCAGCCIT GCATCCCAGG GATGANGCCC ACINGATCAI GGICGATAAG CITTITGAIG IGCIGCIGGA TICGITTITGC CAGIATITTA TICAGGATIT TIGCATCAAT GITCATCAAG GATATIGCNC TAAAAGIGIG CIGIATICAG GAAACCCAIC ICACGICCAG AGACACACAT AGGCICAAAA TAAAGGGAIG GAGGAAGAIC TACCAAGC

SEO ID NO:105: (Length of Sequence = 355 Nucleotides)

GECCTTCCTC AATATGIAGG CGCCACTITI TCTCCCIGIG CCCTCACCIG GTCACCCTC TGIGGGGGAN ATCCCACTGIT CTCTCTGGGGG GTCCAAACTT CCTCTTCTTA GGAGGACACA AGTCAGATTG GATTAGGGCC CACCCCAATG GCCTCATTIT AACTTAATCA CCTCCTTTT GITTGGGCTT TTTAACTTAA TCACCTCTTT AAAGACCTTA TCTCCAACTA AGGITTCATT CTGAGGGTATA CTGCAAGGTTA AGACTTTAAA ACACGAATTT GGAGGGGACG TAATTCAGCC CATAACAATA ACAATAAATGA CATCTTACAA CTTACTGCCA CCACCAAGCT TGCTG

SEO ID NO:106: (Length of Sequence = 355 Nucleotides)

GCATGAGGTC GCCGGGATCG TGGCTGCACG CCACTGCAAG ACCAACATCG TCACAGCTTC CGTGGACGCC ATTAATTTTC
ATGACAAGAT CAGAAAAGGC TGCGTCATCA CCATCTCGGG ACGCATGACC TTCACGAGCA ATAAGTCCAT GGAGATCGAG
GTGTTGGTGG ACGCCGACCC TGTTGTGGAC AGCTCTCAGA AGCGATACCG GGCCGCCAGT GCCTTCTTCA CCTACGTGTC
GCTGAGCCAG GAAGGCAGGT CGCTGCCTGT GCCCCAGNTG GTGCCCGAGA CCGAGGACGA GAAGAAGCGC TTTTAGGAAG
GCAAAAGGGCG GTACCTGCAG ATGAAGGCGA GGGAC

SEO ID NO:107: (Length of Sequence = 273 Nucleotides)

GIGICICITI TAAAGAAAC ATACITIATI TIGGICIAAA TIGIGAAAAT ACCCAAAACA TITGATAGAA ATIGAACICI GICAACAGIG TIATITATAC TAAGATCAGG ACAGITCCII GAGATCATAC IGITTATTA CIAAGITIGG CCITIGITIT ACAAATGIAA IGITCATATI TATITGAATI TIAAGATIGG TIAAATGITA ATGAAAAGCA ATCCAATIGI TANITITITAG TAGIGCCIIT TCICIGIATG CCITAATTIT ATT

SEQ ID NO:108: (Length of Sequence = 359 Nucleotides)

ATTITATITO CITACATOGA AGAAAATGIT AAAGAGTATO TGCAGACACA TTGGGAAGAA GAGGAGTGCC AGCAGGATGI
CAGICITITG AGGAAACAGG CTGAAGAGGA CGCCCACCTG GATGGGGCTG TTCCTATCCC TGCAGCATCI GGGAATGGAG
TGGATGATCI GCAACAGATG ATCCAGGCCG TGGTAGATAA TGTGTGCTGG CAGATGTCCC TGGNTCGAAA GACCACTGCA
CTCAAACAGC TGCAGGGCCA CATGTGGAGG GCGCATTCA CAGCTGGGCG CATGAAAGCA GAGTTCTTTG CAGATGTAGT
TCCAGCAGTC AGGTAAGTGG AGAGAGGCCG GGATGAAGG

SEO ID NO:109: (Length of Sequence = 360 Nucleotides)

TITIATINAAAG CAGITAAACT TAGCATTAAA TAACACTCTT TAAATGGTAC ACCTATGAAG CAAGAGTTAA ATATAAACCC AGTCTAATCC TGTACACTTG TGATTAATTG TGACAATCTT AAGTTGCTCA CTTCTTTCCC ATTTACCAAT TCAGAGAAAG CCCGTTTCCT GTTTTCTCCT CACCACTTTG CCTTGGCATC ACACCAACCC TGCCTCGGGC TTCAGCTGCA GATCCTCCCC AGCCCCTCCT CCCAGCTGGG CTGACTCCAG TCCCAGCCCC AGTCTCCACC AACTGAGCAG CGTACGCAGG GTTGTGTCTG GCTTCCAGCA TCTACCAACC CTTCAGAGCA ACTT

SEO ID NO:110: (Length of Sequence = 364 Nucleotides)

SEO ID NO:111: (Length of Sequence =455 Nucleotides)

SEO ID NO:112: (Length f Sequence = 398 Nucleotides)

CTGATCTGAC AGGAGGTGTA GGTCAGGCAG TAATGGAAGT SATGGGGAAC AGCTGTAAAT ACAGATAAAG CTTTACTCAC
TCGCCCACCC ACTGCTCATC TCCTGCTGTA CTGCCCAGTT CCTAACAGAC AGCAGACAGC TACTGGTCTG TSGCCCAAGG
GTTGGGGACC CCTGACATAG ACTAAACAAT TCACAATGTT TATATTAAAC AACTTATTCC AAGTTTCCAT TTTAGACTCT
GGAACATCTG ACATGGTGAA TCCACAGGTA GTAAATSGGA AGGGAGATAA CAGACAACTT GACGGCCGTG GAAGACGCAC
TGGGGGGGCA CTGGTGACGG GTCTCGGGAC AGACTTCACA TCTCCAGACT GGCACAGTGG GCTCACACCT GCCTCCCA

SEO ID NO:113: (Length of Sequence = 444 Nucleotides)

ATCAGTGTCA GTGTCTAACA GAAGGGTCTG TTAAGGATGC TTCTGATTTA ACCAAAAGAT TAAGCTTCAG AAACAATCTA
ACATACTCAA AGGAGCACCA AATTATCAAC CGGCTACAAG GATGCAAAGG ACCTAAACAA CAGATGTCAA AGGACTTGTA
AAAACTGGAG CCAGCAACCA TTCCACTTGA AGGAATCCAT CTCAGGGAAA TGCTGGAATC CACACACAAA AGCAGGTGTG
CAAATAATCA CTGCAGCACG CCTTCTAATA GTGAACAACA GAGGCAATCC AAATATCCTT CAACAGGGAA CTGAGTAAAT
ACCAACTATG GGCATATCCA CATAAGGCTC TCTGCAGTCA TTAAAAAGGA TTGCACTTAC ATGCATGTCT GCCATGGAGG
TCTTTCAGGC CAATGGTTCC ACTCGGAAGG GCAACCACCA ATTA

SEO ID NO:114: (Length of Sequence = 472 Nucleotides)

TOGGGCCCCA ACGGAGACCT GGGGATGCCG GTGGAGGCGG GAGCGGAAGG CGAGGAGGAC GGCTTCGGGG AAGCAGAATA
CGCTGCCATC AACTCCATGC TGGACCAGAT CAACTCCTGT CTGGACCACC TGGAGGAGAA GAATGACCAC CTCCACGACC
GCCTCCAGGA GCTGCTGGAG TCCAACCGGC AGACACGCCT GGAGTTCCAG CAGCAGCTCG GGGAGGCCCC CAGTGATGCC
AGCCCCTAGG CTCCAAGGAG CCCCAACCGG GACCCAACCC TGCCTCCCTG GGGCTAAGCT CTGGCCTGGG GCACTCACCC
CCTGGCTTAG ACAACTTCTC AAGGGCTTGG CCTTCAGGGG ACCCTTGTGG GTCTTGCCTT GCTGGGGCCA CCTTTTCTTG
CTTGGGGCTT. CCCCTTTGGC CTACCTTGGG GCCAAGCCCC TACCAACTTT GGATTGCCTT CTTGGGGGCC AA

SEO ID NO:115: (Length of Sequence = 293 Nucleotides)

CINGGGGCCA TGTGGCTGAT TTCCATCACC TTCCTTCCAT TKGCTACGGC GACATGGTGC CCCACACCTA CTGCGGGAAG
GGTGTGTGCC TKCTCACTGG CATCATGAGA GCTGGCTTTA CCGCGCTCGT GGTGGCTGTG GTRGCTCRCA AGCTGGAGCT
CACCAAGGCT GAGAAGCACG TGCACAACTT CATGATTGAC ACTCAGCTCA CCAAGCGGGT AAAAAACGAG GCTGCTAACG
TTCTCAGGGA GACGTTGGCT CATCTACAAA CATACCAGAG CTGGTGAAAG AAG

SEO ID NO:116: (Length of Sequence = 448 Nucleotides)

TITIGAAAATT TAGAGGATAT TIATITICTCA GGAAGGTGCA CAACAGCTGG CAGGCACTGC TITICCCTGCT CTAGGGGATT
CCTCTCTCCT TITICCAAGAA ATCCCCTCTC TICTITAGAAG TGCCCATGGG AGGCTGGGAT GTGAAAAGAA ACCATACACA
ACACTCCAGA GCCTTAAAAA AATAAAGCAA CAACCTCCTC CACACGAATA CACTTACAAA ATAAATAGAC GGATAAAAGA
GAGGCCACGT GCCTCCCATC CCGGCTGTAG GGCTGCTTGG GGATAGTGGG GCTGGGTGGC TCGGTCCCAC TTCTCCCAGC
CAGGATGATC CAAAGGCTAA ATGGGATGGA AGGGCCCTGG CTTTCAGAGA GAGGGTGGGG CAGGCCTCTC CTGGTACTCA
GCAGGGAGGA CACTGGGGCA CGCGTAGGGG TCCAAGGGCC ACTTAATA

SEO ID NO:117: (Length of Sequence = 551 Nucleotides)

TACCAATTAA CCCATCATTG CTTTTAAACA ACCATCTGAA GGAGCAGAGA GGCAGGGTAG AAGACAGAAG GGGGTCTATG
TGGGTACTAA AGATGTTTCT GTTTTGTAAT ATTGTGTGTG TGTGGGTTTA TGGTTTGCTT AAGGGATCAA AACCTGGAAA
AAATGGGATT CCAGGAATGG CTCTGTTATT TTTGCTGGGT TCCAGCTTGT AATGCCTACT GCCTTGGTTC A

SEQ ID NO:118: (Length of Sequence = 426 Nucleotides)

CCCCACCCA AAATCAAAAC TEAAGGTAGT GTCAGTGTAT ATATGGAGTC CCTTGTGCTG AAAGTCAAAG CAGCTTCATT
TTGGGGCCTC AAGAGCTCCA GCTCTGGGCT CTTCACCTCT AAGCCCATGG GCAGTGCCCG CCCAGTGGTG TGTATAGATC
GGAGGCTGAG GGCCTCACCC TTAGCTGAGC TGTCGCGTGC TGGGGAGCCT GTGCAGGAGG GTACAAGTAG GAAAGTGCCA
TCTGCATGGG AAGAAAAATG CAGCGTCCTT GGTAGTGCGG ATGGGGTCCA GGAGACCCAG GGAGCTTGCC CAGAGGGACC
TGAGTGGCAT TCCTGTAGGA AAGCAGCCCA GATCTTGGGG CCGTAACGGA TGTTCTGGAA GTTTTGACTT TGAACCACCA
GGTCCCCATTG TTAACAAGCT TCTTGA

SEO ID NO:119: (Length of Sequence = 434 Nucleotides)

TTTTTCGGTT AAAAAGGCCC AAAACTTTAT TTAGTTTTCA GGGAAATATA AGATGCATGT AAACATAAAA TACAAAACAA AACCCAAATC TTACAGTCTA GAAGCATGCC AAGACAGAGC ATTTTCTGCA GACCAAAGAG TCCCGTCAAA GTGATAAAGG ACACCTGGAA AGTGGCAGGC CAAGGGGCTG GTCCCTTCCC CAAGGGCACT GCATTTTTGT GATGAGATTA AAAACAAACC AACTCCACTA TTAAAAATGC TAGAAACATG GGATAGTTTA GCACCACCAT TGATTCTGGC AAATATTTCA GCACTCACAT CGACTGCACT GAGTTTAATG TCCTTTCTCC AGTTTCTCTG CTGAGGAGGG AAGGAGGGAA ACCTGGGCGG AAGGGGCTCC TCCTGACCCC ACAGGGCCAC TAGGAGCTTG GAGG

SEO ID NO:120: (Length of Sequence = 276 Nucleotides)

AGGAAGTGTT AGCAAATGCT ACCATGTGGA ACACTCAACT TTATTTGCTT TATTTATATA TTTAACAATT CTAAAGTATT
TACTTCTTGC TTTGACAAAA AATGAAAAAT ATAGGGGCAC TGACTGACTC CTCTTTAGGA GAAAAGGGTT ATATGTACAG
CTATGGAGAG TTACGGTTCC CCCTTTAACA AAGGCAAATA TTAATAAAAA AGGGCTTCAT CGGTCAAAAA AGGGCTAAGA
GCTGCAAGCA TTTATTCACA CTGTACATCG GGCCCC

SEQ ID NO:121: (Length of Sequence = 554 Nucleotides)

ATTICITICC THATCATAT CTGATGCTGG GATGTGGGTA ACCCCAAACT GAAGGCAGCT GCTAAATCTC AAATGCTAAA
AAAATACTGC AATTITGACA TCAGTGAGTC AGATCAATAC ATCCTCTGGG GCTGATTTTG CTTCACAGTT AGGATGAGCC
ATCTCTTAAG CTGCAGGCTC AAATGGGATT AACTGAACTC TATACCTGGG ATGGGCCATG GACTGAGCTG TCCATGCAGA
AGGACCAGGC TGTCCATGCC TTCCCTGCCC TTTTACTCAC CACTGCACAG CAGCCCCAGT GGGCCTACTG CACATGTCTA
GGAGAAATCA CTCTAAGAAA ACCAACAGGA ACAGGCTTTA GGCAACAAGA GACGTCTCAC TGCATCTCCT CCCACGTCAG
AACTTGAGTA CTGGGTCTTT GCAGCTCAGA GCATTCCTCC CTTCCCTTTC CTGCCCGAAA GGCCTGCCTT TTCCTGAGAC
ATATGGCACT CCATGCTGCA AGTTTCAAGC AGATGCAGGT TCTTATGGGG CTTTTTGCTC AAAGAGCTTT GGTT

SEO ID NO:122: (Length of Sequence = 238 Nucleotides)

CACCIAAGCA GGIAGACATC CGCAAAGTCA GATGCITTCC AACATGACAC CTGAACATCT TCCTTTATGC AACACCCAAA CATCITGGCA TCCCCACCCC AGGAAGTGCG GGGAGGAGGT TATGATCCCT GGGCGCTTCG GCAGAATGGA GAGCTGAGGT GTCCCTCCCC TGCTAGTCAC CTACCAGGTG TCTGAGCAGC TGCATGCTCC CTGGCTCAAG TGGGCACTGT ACCITTTG

SEO ID NO:123: (Length of Sequence = 244 Nucleotides)

ATCCAGGCTT TCATTTCTAG CCAACCCTCA AACACCACCA ACTACAAAGA AAATTTAAAA GTCTAATTTG TAACCTTCAG
ATAAGTATAA ATTAGTTTTT TCTAGGCTTT CATTATTTGG CTTCTTATAC AATCTATCTT GTAAAGTACA TTCCTCTAAA
TTTACATTAT CTAAAATTAA GGCTAAGCAT TATTTAAATC ANTTAATCAT ACAATATTTT ATGGCAATAT GCACATATTT
ATAA

SEO ID NO:124: (Length of Sequence = 330 Nucleotides)

CTCAGCGTAT CATAGGCGTG CTCACCCTCC TCCCCACGCT CCCCCCCCC AGGCAGGTGG TGTAGGATAG AGTGGTGCAT
GAAAGGGGGG AAGCCCGAGG GGCCCCTGG GAAGGGTGCT GCCCCGTAAA GGGCATCCCA CTGCCACTGT GCCTCANCTG
CCGCTTTCTG CTTCAGCTCA GCCAGTCGCC GCCGCTGCTC TTCAATCACT TGTTGTCCCT TCTGCTGCAG AGCTAGTTGG
CCGCTTTTGT TCGATGTCCT GCAGTGTGCC TGCCAGGTTG CAAGGAAGGC TGCCCGGTGC CATTCTGGGG GTGAGTAGGA
GCCCCTTTTT

SEO ID NO:125: (Length of Sequence = 281 Nucleotides)

CCTCTCTCCC TTCGGTTCTC CATTIACCGA GCCACAGIAT TTCTTAAAGC TCGTTGGCAG CCTGCACCCT GCTTATTCTT
GGGAGACACG AGTTTGCATC CIATTACAAC CCATAGTTTT TGCATAACCA TGGTGAGAGG AACCATCCTT CCCAATCCCA
ACCTCAACCA AAGCTTAGAA AAAGTGCCAT CNTTAACCTT TCAGAATCAC TCATAAGTAA ATCCTATAGC AGTCTCTGCT
AATGCAAATT TCAATGTGTG CCCGCTTATT AGGTGACTTT T

SEO ID NO:126: (Length of Sequence = 266 Nucleotides)

CTITIAATGA TGTGGTTCTG GTGGGATTTA TAAAGGGAGA TGGACCCCTG GNAAGATGCT TTCCTMAACC ACAACCCACA CATTGGGTCA CCATTTCCTC TTCCTCCTCC TTCTGTGGGT GGCCGGAGAC CTGTAGGACC TTCCCTCCCT TTAGGGTTCT GTAAGGCCCC TINICAGTCC TCAGAGTCCA TTCTTCTCTT GTGCTGAGGG CCTGCAGTGG GGACCATATA CTTCTGGTGC TCTTAGTTTG CTGTCGCGTC TGTTTT

<u>SEO ID NO:127:</u> (Length of Sequence = 435 Nucleotides)

GECTIGITET ATTEATITIE TAGTICCEAG AAAAGGAATG AACCIGACT ATGGCAATIC ACCIGACGT GIGATAATITI
AGITIGCIAT GAGTITICAC TETTAGGIAA AACCIAGITA TECTAATIAA TAATTAGITA TEGATGATAT AGIAATITITI
TITTITITIG ACTGCGICIC ACTGICATIC GGGCIGGAGT ACAGIGGCIG ATCACAGITC GGIGCAGCCT CGACCICCCT
GGGCICAGIG ATTCTCCTGC CICAGCITCC CAAGIGGCIG GGGATTATGG GCATGCACCA TCAATGICIG GCTAATGITT
GGIGIGITTT TITATAAAGC CAAGGGTTTT GCCCATGNIT CAAGACCCCG GGGCIGGICC TIGAACCICT TIGGGGCTTC
AGGCAAGTCC TCCCACCITC GGGCCTTCCC AAAGT

SEO ID NO:128: (Length of Sequence = 471 Nucleotides)

TTCCCTTCCC AAGGACTCGA CCTGAGAACC GCCATGTACT CGGAGATCCA GAGGGAGCGG GCAGACATTG GGGGCCTGAT
GGCCCGGCCA GAATACAGAG AGTGGAATCC GGAGCTCATC AAGCCCAAGA AGCTGCTGAA CCCCGTGAAG GCCTCTCGGA
GTCACCAGGA GCTCCACCGG GAGCTGCTCA TGAACCACAG AAGGGGCCTT GGTGTGGACA GCAAGCCAGA GCTGCAGCGT
GTCCTAGAGC ACCGCCGGGG GAACCAGCTC ATCAAGAAGA AGAAGGAGGA GCTGGAAGCC AAAGCGGCTG CAGTGCCCCT
TTGAGCAGGA GCTGCTGAGA CGCCAGCAGA GCCTGAACCA GCTGGAAAAA CCACCAGAGA AGGAAGAGGT TCACGCCCCC
GAGTTTATTA AGTCAAGGGA AACCTTCGGA GATTTCCACA CTGACCAGCG AGAGAGAGAG CTTTAGGGCC A

SEO ID NO:129: (Length of Sequence = 186 Nucleotides)

GCCTTIAACA TCCTCTGCCA ATRACTGGCC TCAAATCACC AGTGGAACCT TTTCAAAAAA TACACCATTG GCTCTATGTA
GTTCTACTGA TCTRAAATAT CCACGTGTGG GCCAGGAGCA CTGGCTCATG CCTGTAATCC CAGCATCTTG GGAGAGCGAG
GAAGGAGGAT CATTTRAGCC CAGGAG

SEO ID NO:130: (Length of Sequence = 307 Nucleotides)

ATAAAATACT TAGGAATATA CCTAACCAAG AAGGTGAAAA ACCTCTCCAA GGAAAACTAT GAAACACTGC TGAAAGAAAT CATAGACTAC ACAAATACAT TTCATGCTCA AGGATGGGTA GAATCAATAT TGTGAAAATG GCCATACTGC CAAAAGGGAT CTWCAAATTC AACGGTATCC CCATYAAATA CCACCATCMT TCTTTACAGG NITCGGAAAA GGAATTCTAA AATTCATATG GGACCCAAGA CGGGCCGC ATAGCCCATG GCCGCTTAG SENWAAGGGA CAAATCTGGG AGGCCTT

SEO ID NO:131: (Length of Sequence = 184 % sotides)

CCAGGITGGA TGGAGTGCAA TGGCACGATC TCGGCTCACT TAXCCTCCC AGGITCAAGC AATTATCCTG TCTCAGCCTC
CTGAGTAGCC GGGATTACAG GCACGTGCCA CCACACCCAG CCAATTITTG TATTTITAGT AGAGACGGGG TTTCACCGTG
TTAGCCAGGA TGGTCTCAAT CTCC

SEO ID NO:132: (Length of Sequence = 270 Nucleotides)

GENEGAGGGE GTOGAGGGEC AGGAGETATT CTACACGCCC GAAATGGETG ACCCCAAGTE AGAACTMITE GMENAGACAG CCAGGAGCAT TGAGAGCACC CTGGACGACC TCTTCCGGAA TTCAGACGTC AAGAAGGATT TCCGGAGTGT CCGCTTGCGG GACCTGGGGC CCGGCAAATC CTTCCCGNNNC ATTGTGGATG TCCACTTTAA CCCCACCACA GCCTTCAGGG CACCCGACGT GGCCCGGGCC CTGCTCCGGT AGATCCAGGT

SEO ID NO:133: (Length of Sequence = 529 Nucleotides)

CITGCAGTAC ATACCATTGT TATTACTGAT AGCTTTATAA ATCTGCCAAA TAACATAGAA TGTAGCCTCA AAAGGATGGT
CGAGGGTTCG CAATCTTTCT TTCTCCACCC AGTGGTGTGG AGCAACTCTG TGCCTTAAAG AGGGCACCAT GGAAAGAAAC
AAAAAGGAAT CTCTTTCAAA ATGCTGGAAA TTAGGCTTAG CTCACTACTT TCAGGATAAA GACAACTGCA TCTAATTAAG
TCCACTCCAC ATTTCTTTGG ACTCTAAGTA TTCTGCACCT GAAGGCTAAA TTGAACTGGC TCAGCCCTAT CTTTTTTGCC
ACATCTTTAA TTACAAATCT ATTTCTTCTT CCTTTCATTT ACTTCTCTTC TCTTAAGTAA GAAATGTGGG AAATGAGACT
GGCAGTTTGG TTTGTTTGCA TGTGGGTGC CATTAGGCGT CTCATCCTAT GGCCCTTTTT GGAAATGTTG CCTTCCTACT
ACACACCTGG GAGGTTTCCC CAAGGCTCAA CCTTTTTGCT TCAGGTAAA

SEQ ID NO:134: (Length of Sequence = 437 Nucleotides)

GACGGTGGCG ACGCGTGCAC CGGGGATGTG TCCTGCCACC AGAGGAGGTG TGCGTGGCGG GGAGCAGAGG GGCTTTGTTT
CCCAGGTGAA GGTGCGGCTT CTTCACTCTT AGAGGTGCGT GTGTGGGGTGG GGGTGCTTGC TGTTGAGGTT TATGCCTGTA
ACTGACAGCT GTCCCCCAAG CCATGCTGGC AGTGTGTAGG TGTCGTGCCG GCCACCGCAG AGGAATCCTC TGGGCTTCTG
TGGTTCAAGT GGGGCCCAGC GCAGAGCTCC ATGAGTTGCT GAGCAGCCAG CCCTTCAGCA TCTCCTGGGT TTTGGCAGCA
GGAGGGGTCC CCTTGTGCAA TTCAGGGGGC CGTGGGGGCACTC GTAGCAAGGT AAAGGAGCCC CTGCTCAGGC
CCTTGTTTGC TCCCCTTTCT TGCAAGAGGG GTAGACG

SEO ID NO:135: (Length of Sequence = 534 Nucleotides)

GECATTGITC TOGINGGIGI GICACGCICC CAGAAGACIG AATTIATOGI AGGATCACIC GCAAGGCCIT GIGAMGCATI CITACCIAAA ACAAAAGAAA TATCAGGGAC TITIGITGAC TATTIACAAC TCAGITTIAC ATTIAAATTC AGGCAGIGIT AATATGCCAA GGIAGGGAAT GIGCCITITI CAGAGTIGGC CAGGAGCTCU TGGCIGGGAC ACGGAGAGGC AGGIGTGGCG TAAGGCCTCA CTCCCGGCTG TGAAGGTCTC TGATCACACA GAAGCAGCCC TGCCCAGGCT GGGTCATTTG CTGTCCGCTT
TTCTCTGTGA CCACAAGCAG CCCTGAACAA CCAGTATGTG TCTTCTTTCT CCAGATAGTG AAAAAGGGTG TCCAGATAAA
CCCACCTAAG TGAAATGGGC CATCCTCTAA ACTGGGGTAC CTCACTGCAC AGGTTCTAGG TAGGCTTTCC ACTTAATCTA
ACTTGAGGCC TACAGGTACC CTGTAAAGTT AGTGGGGCTT GTCCTTGATT GTGG

SEO ID NO:136: (Length of Sequence =279 Nucleotides)

CAGTITIGAC AAAGIAGCAT AGIGACITIN TITCTIACANT GACTITICGGA GAAGIINGCA GITTICTIGGCA AAGIGACGCT
GGGCTGTITIG AAAAAGGCAA GCTTAGCCTA GGCTGCCATC TTAAAACATT TOGAGGCTGT AGCTTCCTCA GGATCCTTTG
CCTGTGGTCT GGTGGCCGGC AGIGCCCCGT CTAACAGCTT TTAACTCTGC ACTTAGTGCC TGAGCACCTA TGGCTGTGAG
AGATGCTAGA TACAGAACCC TGTCCTGTAC CACGTGGGG

SEO ID NO:137: (Length of Sequence = 518 Nucleotides)

CAAATATTIA ATGGAGATCI TCCITGITGG TCTGITATAT GICTATCCGI TTCTGGGTGG THTAGGAGAA TCTGTACTAT
TTCAGCATGI CCTCCTCCAG CAGCAAAATG AAGAGGAGAA CTAAGITGIC CATTIAAAAG GITTGGATTG CACTITCCTT
TCTCTAACAA TATGGGAGTG GCCTCAACTI TTCCATACCA GCATGCATAA TGAATGGGTG CCCAGTGGTC ACTATCTAAC
TGGTTGACTG AAAATCITTC ACTGAGAAGA CGGCTTAGTA ATTCTGAATC TCCTTCACAG GCGCTTCGGT GGAGAGGAAA
ATCATCTACC CACTGTCGTT CCTTGTCTTC TGTGACACTG CTCATGCTTC TCTGCCAGIT TTTCCTGTTT AGGGTATTTG
GATTTTTGAG TAGTCTGGAG CTCCTAGACC CAAGTATGGA TTTATTACCC ACTTATCTAC CCGATTTGTA TACTGAGGAT
CCTATCCAAC AAAGGGTGTA AATCCAGGAT CCGCCTTC

SEQ ID NO:138: (Length of Sequence = 266 Nucleotides)

GATTGCAGGC ATGANICACT GOGCCCAGTO GAGTGGTAAT ATGTIMAAAG GAAACCTITT TOTGAGCAGG TOTCAAAAGA
GAGGTTAAAA TACTGAGTAG ACCATMCTGT AAACAGATGT MCTGTTATYC GGGCTTTCAT ATTCCATTTA TAAAGCACAG
GCAGAGCTCA GAGTAGATTT AAYGTAACTO TGAAGGGCAC TAGGATTTTC AGAATGGTAA ATAAGCATTG GCTTCACCTT
AAATYCAAAT CTGCATTGGG CTTGTA

<u>SEO ID NO:139:</u> (Length of Sequence = 341 Nucleotides)

ACCTOGOTCA COGCTOTGAC CACOGACAGG CAGAGCAAAG GATGOGGGAG TTGCCTCTGC TGCCCATCTA AGGGGACGTA
GGCAGAGAAG CAAAGGCCTC TGCTCTCCCT CCATCCATCC CGGTGTGCTG GCCCCAACGG AACAGGAGTC CTTCAACTAT
TGCCTGCCAG AGACCCAATT TTAGGGACTG TAGTCTGCAT CTGGATGAGC TGGGCTGTAG ATTGAAGTCT CAGAAGCAGG
GAAGGTTGGA AGGGGTAGGG TCCCAGAGCC CATGGAGTTA TTGCTGAGAA GATATGCAGG GGACACATTT CCCAGGGGCA
GAGTAGAAGC CCTGGGCCTT G

SEO ID NO:140: (Length of Sequence = 234 Nucleotides)

GTGAAGGGAG TTGCAGAATC AAATTGCTAC ATAGGCCAAA CAAAAAAGAA GGCTTTTTCA AAAAACATTA AATTCACATG CAGTCTCAGA GACTATTTAG GCAAAGTTCA AGTTAGGAGC TTTTAGGATG TGGGANTAAA ACTTTAATKG GAGGGGAGGG CTTGCTTCTG GAGAAGGAAG AAGCCAGACT TGTTAGACAG TACTCTTAAC TCCTAGCCCA GCCTAGCGTG CCCT

SEO ID NO:141: (Length of Sequence = 354 Nucleotides)

CHACRAGGI TAGCAACTGC AGGAAAACTI TCTTCATTTI CACTGAATTI TAAAGAGAGA MYUUGICTC TATTTCTCAG AGAAACTTAG GIGAAAAGTA AAAGAGAGGC AAAATCTCTT TCCTTCATGA GATACTTTTA TTTTTATCTC TTTCTCTACT CATGTGCTTA ACTGGTGAAA TGATTCTGTA GAAATAGATC CTTCTGATTC TGCATCTCAT TTCCTTATGG CAACTACAAC AGGAGGAATC CAGCTGGAAA TGCCACTAAC CCCACATCCA GCACCTGAGA GAGGAAGCCA GTCGGAGGGC CGTGCTGGGC TCACTCACTC TGGGCCTGCG CACTGGGTT GTGG

SEO ID NO:142: (Length of Sequence = 373 Nucleotides)

GITTITICAA CACTITITIT TIAAGITATI GGGIGCAAAA TCCCAAACCA GGATATGIGI ATGICIGIGI GITTATGITT
TINATITGAC CCTCCCCTCT TICAACCTAC CCCCTITTAT ATCIAATGIA GAAAAAGCGA AATTGAATCI GGAAAGCAAA
CTGTTGTATA TAGTTGCGGT AACAATCATG AAGAGAGAGC CGGGCTGTCC AGTTGTTTTT GAGACAGAGT CTCACTCTGT
TGCCCAGGCT GGAGTGCAGT AGCATGATCT TGGCTCACTG CAACCTCCCC CTCCCTGGGT TTAGGGCGATT CTCCTGCCTC
AGCCCTCCCA AAGTAGCTGG GATTACAGAC CCGTACCACC ACAACTGGGC TAA

SEO ID NO:143: (Length of Sequence = 262 Nucleotides)

COGCACCTOG GCCAGAGGOG GCTGCAGCAG CTGCTMCCTT TTCCCTGCCG CCGCCTCTCC AGTCCCTTTT TTAATTACCA
CTCCAMCTGC TGGGAACGGG CGAGAAAGAG GAGGAGGCGA GAAACTCCCA CCGACCCACA GAGGGAGCAT GATTTCGGCA
ACTTCACCTA TCATTCTGAA ATGGGACCCC AAAATTTTGG AAATCCGGAC GCTAACAGTG GAAAGGCTGT TGGAGCCACT
TGTTACACAG GTGACTACAC TT

SEO ID NO:144: (Length of Sequence = 384 Nucleotides)

GGAAAAGCG GACCCAAACA GTGGTGCTG GGAAATTTT CCCTGTCCCC TTTGGAAGGC TGAGTGGGTG ATGCAGCACA
GGAACAAGGC TTGGACGTCA GAGGTCTCAT CTTCACTGIN ACAAAGCATA AAGGACTTGG GGTTGAGCGT GTGINTGGGC
TCAAGTGACC ATGCAAGTCC TGTCACCTCC TTCCTAAGAC CCCATCCTTC TCCCAAGTCC TCCACAAGAG CTACCTTCTT
CAAAACAATA ACAGAAACAC ATCAAGCTTG GGCGTCACTG AATTCAAGTT CTGATTTCTC CCGTCACCCC AGCAACAGTG
CCCAGTTTGA TTGTGACACT TTGACCCAGC ACTTGGTTTT GAATGTTCTT TTCGGCTTGT ACCG

SEO ID NO:145: (Length of Sequence = 324 Mucleotides)

CTACATGGAA TCATAAGIKT TCCTAAAAAA GGAAGACAGA TITGAAGACA GAGGAGGAAG GTGATGTGAT GATGGAAACA AGGGAAGAAA ACGCAATGIG ATGTGGCCAC GAACCAAGTA ATGAGGACAG CCTACAGAAG CTGGTCAAGG CAAGGAAACA GATTCTCCTC TAAAGTCCCT GGAGAGGGCC TGGCCATGCT GACACCTTGA TTTTKTCCCA GCAGAAACTC ATTTTGGATT TCTGGCCTCC CAGAAAAGTA AGGGGGAAAT GTGCTGTTTT ATGTCAGGTT TKGGGTAATT TGTTTATTGC AGCCATCGGGAAGG

SEO ID NO:146: (Length of Sequence = 355 Nucleotides)

TTTGCCTCCT TCCTTCCTTA TCCAAGCAAG GGTGTGGTGA CAATGACCTG ATCGGGGTTT AACGCCGGCT CTGTCTGCTC
ACCAGACCTG GGGTGCTGAG CTCTGACCAG CCTGGGCAGC CCAACCCCACA GGAACTGCGG TTTCATAGCT GGGTCTTCAG
GAAGGGGTGG AGGCTTTGGG AGTGGCAGCT CCCCGCCTCC CACCACCCCA AGCCAGAGAA TGGGGCAAAC TTGTATGCAT
GGCTTATCTC TAAATTACTA ATCTGCTTCG GACCAGACTC ATCTCTACAG TATAGAGTTA GAGTTATTGC TTCTATGACA
GGTGTTCCAG AAGCCCTGGG TGGCTTTAAA GTCTG

<u>SEO ID NO:147:</u> (Length of Sequence = 337 Nucleotides)

CAGITITCIG AGITCCCGIG TGCIAGACIG GCCAGAAGAG AGGGICIGGG GCCTGGICAC TCGGCCACIC TCTCCTGITT CIGGCCICIT CICCCTICAC TCCCGICCAG TCIGGITTIG AGAGCAGGGG CIGGICIACA GCACCTCAGG GAAGGGAGGA GAGAIACCIG CIGCITCCAT TGCITTICCC TTCCTGGAGI CGATGCCTIT CTAAGGGIIG GAGCIGCTCC TTGCAGGGGC GGGTCAGTTT CCCAGGCCAT GCCGGGGGTG GCCATCTATG CTAGGGCTGG AAGCTGAGGC TGGCCGCCAA CTGTGGGGGCT GGGGTGGGGG TGGGTGG

SEQ ID NO:148: (Length of Sequence = 278 Nucleotides)

GGAATCAGAT GCTCAGGTGT CCAAGCAGGG ATAAGGACAG GCAAAATAAA TAACCCGCCC AACCCCCATC GTCACTCTGC
TGCAACACGA CACAAAGGTT TAAAGATCTG GGCCCAAAGA CTCTGGGGACC CTTCAAGCAA GTCAGGTGGA AGAAGGTTTC
CCCACCCCCC ACCAGGCCTG TTTGTCCCAG GTTGCCCTAG GATGGAGGCA GTTCAGACCC TGGGTCACTG ATGCTTGATA
GGAAGATCTT TGATATCAAT GGCCTAAGCT CTGCTCAT

SEO ID NO:149: (Length of Sequence = 368 Nucleotides)

TITITITIT GITTICAACA AACITTACIA AATAACCCIG GAAAGGCAAT GAACGATCIG ACAATITAAG CICTAATGAT
TIAAAGCICA GCIAGAAGAA AGIGAGGCAT GACATATACI GICAACGGAG GGIGAAGGAG GCAGATITCI GGAAATGCAA
TGATCCCACA CATTIGCIIC AAGGAGAAAC CIGCAGACAT ATTITCAGGI CIIGCIAAGI AACAACIGII TATTIGIAAT
CAATACATIT GGGGAAAGIC TGCIATGIAG CIAAGGICAC TGIGACCACA GACCAACAGA TGGAAAGGAA AAAGGCACIG
GACCAGCAAG GAAAAATACA TCCCCATCCI CAAAAGAATI TIAAGGIG

SEO ID NO:150: (Length of Sequence = 367 Nucleotides)

TTGTGAAATG GGCCTGGGTA GATAAGGAAA AGAACCTCCA AGAGGTTAAG TGATTTGCGG ATTTGCCTAA ATTATACAGA
AGAGTCAGCA CCAGTGCCCA GGCCTTCTGA TTCTTAGTGC AGTAAACACT AAGCACCATC ATTCCATTTC ACCACACTCC
TGTCTTGCTG TTGTCCTCAG CTAAGAAAGC CTACCCCTGA GTTACCCTCT TCCATCTTAG AGCCTTCCTG CTCGCTGTCT
GCCCCCCTGC GATGGGGACT TCTTTGGCCC TTCTCACCCA GCCCAGCCTC TGCCCGTTTT CCTTCCTTT TCCACTGCGG
CTGAGCTCTT TTCTCCTTCC GAGAAGCCTT TCCTTCATCT TTCCTGG

SEO ID NO:151: (Length of Sequence = 366 Nucleotides)

CCCAGCGGGC CGCCTCCCTC CTCTCTCCTC CATAGGIGGG GGTTGTGGGC CTTCTTTTTT TTTTTGTCTT GGAGGGCAGT
TAAACTTCTC CATTTGCCTC TCTCTTCACA CCCAAATGCC AAAGGACACT TTTCCTTTCT TTTTGTGGGTA GTTGCAAAAA
AAAAAAATTC CTATGGGTA CTGCCACTTT TAAATACTTT GTAACTTAAA GGCAAAGTAG TATGTCACTG TTTCTTTTCC
CTGTAGTTTA CTTTTGAGGT TAAACATCTT TCCATGTCTT TATTGGTCAA ATACAGTTCC TYCTTTTGTA CAATGTTAAT
CCTAATATGG ACCATTTTTC CTAATGGGAT TACCGATTTT TTTAAA

SEQ ID NO:152: (Length of Sequence = 269 Nucleotides)

GITATTCTGG CAAGTGCTTT CAGGGCCCTC CAGGGTTTGG CTGGTCACCA TGGAGGGGGG GTTCAGGTGC TGAATTTAGG GACCCCAGCA TCTCACAGGT TTCCCCCTTCC ATCTTTCCCA GTGGCACTGT GTCTGAGCAG GTGTGCCCAG GTGAGGTTGT ATCCACTGTG TCTGAGCAGG TGTGCCCAGG TGAGGTTGTA TCCACTGTGT GTGAGCAGGT GTGGCTGTTG CAGGTGGAAG TGGGGATATN TGGGCACCTG GGTGCCATT

SEO ID NO:153: (Length of Sequence = 260 Nucleotides)

TITICAGGATT TIATTIAAAA TITATIGIAA TGGGGTCCGC GCAAAAGGAA GGGGTGGAGG GTGGGGTACA TGCAGGGGAC ACAGGAACAN GATCCACATG GCCAGGGCCA CAACTTCITC TGTCGTGGGG AAGAGGGATG AAAAGACAAG ACCAGGGCTA NGAGCTGGGG TGGAAGAGGG GAGGGGNAAC ACTGGCTGCA TTCCCCNAAC CCCANGANGC ACCTATAGGC CCTGGACCCA TGGGTCACCC TGGGCCCTAG

SEO ID NO:154: (Length of Sequence = 405 Nucleotides)

TEGRACTITET GAGTGGGGAC CCATGATGTA TEGGTCTCAC CTGACTTGAG GTGAATTTTG GAGTGAAGGG CCCTGAGGTC AGCTCCCAGG TCGGTCGTGC TEGGCCAGGC CTGGTTTTCA CAGGGGCTGA AGGATCCCAG TCCACCTGTG TGCATGTCAG GGCTCGGCCG GGAAGAAGCC AGCAAAGTCC CCCGTGTCCC TTGCTGAGTA TTCTGTCACA GACAAGCCTC CATTAAAGCC ACAGCAGTGC TACCCACCAC ACACACCTTG CTGGCCCGGC CACCACTGCT GGCTTCAGCC CCTTNAGCAG CCCATGCNTT AGCAGACCCT CAGATGTAGG TCAGTGGCCT TANCTGTNTC TATCCATGCT GTTAAACTCC CTGCCTCCAA CTGGGGGTCA CCAGT

SEQ ID NO:155: (Length of Sequence =: 40 Nucleotides)

CCATGATCIT ATTIATIACA TCIAGITITI CITTATACCI CIAAAAAAA GIGCCITITA GATTIACAGC TIGIGCITCT AAAGCAAAGG TIAAAACATC ATGCCCCAAA GGAAAACAAG GIAAAAAGGA AGCIGCCATA TAAGCICITIA AAANITGIAT GITACAAGGI TCIAAAATCI CITCAGCACI GGIIGGITGG TAGATIGIAC GACACTGACA TGGIGCITGG GAGGGICATT TATCIGATGG TIGGAGCAGC ACCATGGGAA AGCIGCCCAG ATGGICTACT GAAGTCCTIG GCIGIGCACA GAATCGCCCC AAGGGCCAGN AATTCATGAG TCCGGGGAAC TITGGNGGIC CITACTCAAT CTCCTTAGTG CTAAAGNITC AGAGTCTCAA

SEO ID NO:156: (Length of Sequence = 443 Nucleotides)

GICCICIGGA TIGCITCGIT GGITGCGAAC TITAAGAATG GCAAACTGIG ATTGGNICCG ATTAAGACAA GCITTGTAGT
TITCITCGIG TAAACACCAA ATCCCGCCTG GGCCATGAGG TAGCAGAAGT GGCCGCATC CAAGAGGCCC CITGAAGCCA
GAGIGICGCC CATGGTAGCC ATCGICCTGG ACTCGACGIC CATGITGITG TICAAGTTGG ACAAGACCAT GGCGAGGTGC
GGCCTCCAAT CICCCCATTT CICGICTCCA CAGCACGIGG ACGCGGCAGG CATCCGTCCG GACATGAGCT GGTAGACTGI
CTTCAGAGGG TCGTTGATTK GGGAGGCTTT TTAGCAAACC TKGGTCATGA CTCGGGCGIG TGTCCGGCTG TTCCATCITA
CTTGCAAGTA GCAGAGCGIG ACCCCACAAG GCCATTCTTA ATT

SEO ID NO:157: (Length of Sequence = 383 Nucleotides)

ATTGGAAAGG GTTTTAAACG GAGICGGAAC CIGAGTAGAT TICCAAATTT TACAGCCAGG ACTACAGAAG TGCATCATTC
TAGAATGTGT AGACCTGAGT AGCTTATACA CTACAGAGCA CTTTGCTTAT TIGAAAGTAA TICAGCAACA GGTCACTTTG
GGATATAACC TGAACCTTTT TITGGAGTGG GGTGGGTAGA CTACAGTAGA CACAAGGGCT GGACATGCAG ATGCTTAGGG
GATTAGCGTT TITCATAATT TGTTCTGTTT GTCAGTTCAT TCCTGTGTGT TCTTACCTCT ACAAAGGTAC ATTACACATT
TTARGTTTTT TAGTGACCTT TAACCATGTT ACTTGAAGCA TTTTGGAATA TAAAGCTATT TTA

SEQ ID NO:158: (Length of Sequence = 241 Nucleotides)

TEGTISTIGIG CTCAGCTIGCA GCGGCASGTA AGIGGGTSTC CAGGGGAGTIG GACAAGCAAT TCTCCTGTCA TTTGCAACTT
TCTTCAGGAA CTCAGATAAA GAACACTTGG ATAACGATGA TCCCTGTAGA GGGATTTCAT CTGTACCATC ACACATGGAA
GAGGAGTTTC TAGGTCAGGA AAGGCAGCTN CTAAGCTAAA GGTTTCTTGG TCCCTTNGTC CTGGCATGCC TTAAGGAGGG
G

SEO ID NO:159: (Length of Sequence = 224 Nucleotides)

CTGTCAGTAA TGGCTCACTA AAGGGCCAGC AGTTTAAATT ACACAGGTTG CACTAAAAGC TGCAGCTTTG GCCAGGCAAG GTGGATCACG CCTATAATCC CAACACTTTG GGAGGCCGAG GCGGGCAAAT CACCTGAGGT CAGGAGTTCA AGACCAGCCT GGCCAATATG GTGAAACCTA AGCCTCTACT AAAATTACAG AAATTAGCCG GTCGTGGTGG CACA

SEO ID NO:160: (Length of Sequence = 377 Nucleotides)

in te

GAAGCCIGAG GCGGCCGGAT CACGACGITA GGAGATCGAG ACCATCCTCG CTAACACAGT GAAACCCTGT CTGTACTAAA
GATACAGAAA ACTGGCCGGG CGTGGTGGTG GGTGCCTGTA GTCCCAGCTA CTTGGGAACT CGGGAGGCTG AGGCAGGAGA
ATGACCTGAA CCCGGGAGGC GGAGCTTGCA GTGAGCAGAG ATTGCGCCAT TGCACTCCAG CCTGGGCGAC AGAGTAAGAC
TGTCTCCAAA AAAAAAAAAA ATAATAATCA AAGCTCTTGG ATTTATAGTT TGGTCCCCAG CCTTGTTTTG ATCTTTCCTT
TATCCTGTTT TATTGCCATT TACCACGTCC TTTTGGAAAC ATCCCTTTCA ACTGCTG

SEO ID NO:161: (Length of Sequence = 273 Nucleotides)

GCAECGGCGC CGGGCGAGGA GGCGGCAGGG GCGAGGAGGG GGCGGCGGGT GGCGACCCGC AGGAGGCCAA GCCCCAGGAG GCCGCTGTCG CGCCAGAGAA GCCGCCCGCC AGCGACGAGA CCAAGGCCGC CGAGGAGCCC AGCAAGGTGG AGGAGAAAAA GGCCGAGGAG GCCGTGGCCA GCTCCGCGT GCTAGGCCCC CTTCGCGCGG GCCCGGCGCG CCCCCGGAGC AAGGAGGCAG CCCCCGCGGA GGAGCCCGCG GNCGCCGCAG ACT

SEO ID NO:162: (Length of Sequence = 286 Nucleotides)

TTTTGGTCAA ATAAATCAGA GTACTACAAT CATCAAACAT CTGATTCATT TAACATGTGA GCATCTATAC CTGCCCATTT
GTGTGAATAT TCAGTATATA TCTCATACCT ATTCTCATGC CTTCATTTAT TGTGGTTATG GCTGTAGATA TGGAAAAAAC
AGTAGCTGAG ACATTTTTAT TATGAACTAT ATTATACCTT AATCAATCAG TCAGAAAATG CTTAGGAAGA AGAAATGCAT
GATTGTAAAT GCATGATTTC AACATGCTAC CCGGCCAACA AAGTTG

SEO ID NO:163: (Length of Sequence = 342 Nucleotides)

TGCCCAAGGA AGACAGAACA TGGAGAACCG TCAAGGCAGG AACCCCACAG ACTGTCCTT CCAGCCCACA CTCTGCCACC
TCCTGGCCCT GTCCCAATTC TGAGCCAAGG CCTCCCCGAG GCAGAAGTTG CCTGGTCCTC TGTCCCCACA GTGACCTGAC
TGGGGGTGAG GGAGAAGGAG GAGAGAGCCC ATGTGTGGTG TGTGTGCCCC TGAGAACTTC GTGGTGACTG CCTTTGGGAG
CCCGCAAGTG GCCAGAGGCA GGGGTAGCTG AGTTCCTGGG AGACCCCTTT TTTTCCCCCA RGTTCCCCAG AGGGCAACGC
CATCAGTAGC AGTGTGGTGT TT

SEQ ID NO:164: (Length of Sequence = 392 Nucleotides)

ATTACCOGGE COCCECTOC CTAAAACAGA TOTACGGACC TTAACCGACG CCATGOTGAG GOTCATTOCA TOCCTGCRGA CGTATGCAGA GCCCTCACT GCTGCCATGG TGGAGTTCTA CACCATGTTA GGAGGAATTC ACCCAGGATA CACCACCTCA CTATATCTAT TCACCCCGTG AAATGACTAG GTGGGTGAGA GGCATCTTTG AAGCGCTGAG ACCTCTGCAG ACCCTGCCTG TTGAAGGCCT CCTTCGGATT TGGGCACATG AAGCTCTGCG TCTCTTCCCA GATAGACTCG TAGGGGATGA GGAGAGGCGT TGGGGACTGAA TGAGAAGATC GACACGGTTG CTCTTGAAGG CACTTTCCCT AACCTTCGGC AGAGAGGAGG GC

<u>SEO ID NO:165:</u> (Length of Sequence = 406 Nucleotides)

GITATAATTA TCTTGTTTTA TTATTTATTG TITATCTCTT ACTGTGTATA ATGTAGAAAT TAAACTTTAC CATAGGTATA
TACATATTGG AAAAAGCATC TTATATACAG GGTTTGTTAC TATCTGTGGT TTCAGGCATC CACTGGGGGT CTTGGAACAT
ATCCCTTGCA GATAAGAGGG AACTGCTGTA TCCATAGAAT AAAAACACCC CATCTTGAAG ATAGGAGGTT CTGTAAATTG
GGATGGGGTC AGGGAATCTG AATTTTAAAA GTTTCCCATG TGATTTGATG CCCAGCCAAG GGCTGGGGAC CACTGTCTTG
AAATATAATG CTGAGGAAGA TACTGTCTTT GGATTTTCCT GGTAATTCCG AGTGCAAATT CTCAGGCTGG AACCTTATGG
GCCTTG

SEO ID NO:166: (Length of Sequence = 453 Nucleotides)

GAAAACTTIG CCATGGGICA GITTIATIGG AAGITCATIT TCCIGAATGI TIGGAAGAAA GICIAGIGAC TCAGGATAGC
ATTICIAATT TCACAGAGIT ATTITICCGI TATGAAACAC AGAITGCCIT TGAGGICICC TGITICIACT ACTGCCCCIC
ACTITIATGI GGGCCICCIC TITCCITIGI TICIGGAGAA CCITITCCIG TICAATTCIG TITTAATTIT CAGCAGITIT
TTITCIGIGI GAGIGAGGCI GITICCIAGC AGGGAGGICI GGITGGICAT TITCAAGITC ATCAGGGCIT CATCAGGGCI
TGICCACTIC AACCCITACG CIATAGGICC CINIGCACCA TCIGCANICI TCAAAATGIG CCCACIGGIT CGITCCCATG
GANGGCTTGI TGGIAATTIG GGCITITAGG GGGGGCCATG GAAGGAGCAA ATC

SEO ID NO:167: (Length of Sequence = 285 Nucleotides)

TITIACTOTTA AAACTGITAC AACAGAATCA TGGACTGACA CAGGTAATGG CTGAGCCATA AGCAAATCGA GAAGTACAGA
AATGICCCAC CCCAAACAGC TGCGGAGTAC ACATCACACA GGGCCTCTGG TCCCGGCCTT CTCAGGTGCT CTGGAGTGGA
GGATCCTTTG AGGGAACTCT GACCACTCCT GITGTCTACC TAGAGAGCAC GCCACTTGGG CCACCTACCC CCAACCTTTG
GCCAAAGGAG TGAAAGGACC TGGAACCTGT CGTCAACCTC AGCAT

SEO ID NO:168: (Length of Sequence = 327 Nucleotides)

CTAGAGGGCA CTCTGTATAC CCGTCAGCTC CTGGAGCCAT TCATTCTATG CTGGGCAGAC AGGCTGTGAG AGGACATGGG
GGACGGTGGA AAGGNTCCAA AGACGAAGCT GINGTTTATC CTTGTTGGTT TTACACAGGG AATGATGAAA CATTGAAGGG
GTTTAATAAG CTTTTCCTAA AACATTTTCC CCCTAAACAG GCTGGCACTA TGTCGAAGCT GCCCAAATTT GAGATTGATT
TACCAGCTGC GNCTAAGTCA ACTAAACCCA NGCCTTTCCG AAAGAGACAT CGCAANTGGC TTACCCAANG TANTGTCCCG
TTTTCAG

SEO ID NO:169: (Length of Sequence = 346 Nucleotides)

GGIGCTATGG AGAGCCGGCC GTCCTCCAGG GGIGAGCTGG GGAGGCTTCT GCGGTTCTGG AGTCCCGGCG ATGGCGCCAG

TTCCCCCAGCA AACCCCCTCC AGAGCTGCCC CCGGATGCAC AGACAAGGAG GGGGCTTGGG AGTGACTTGA GGCTGTGACG

GGRTCGCCCT CGGTGTGGGC AAGTGAGTCC TCTGTGGCCA AGAGGTCAGA GTCGTCCCTG AGGCTGAGTC GAACACAGAC

CCGTGGCCCCT CATAAAATTA AACATAAAAG CACAAAAATG GGCGCAACCA GACAGCATTG GCTTTCAGAC AGGCAGGGAC

ACGGGGCCC CITCGIGITG ACCIGT

SEO ID NO:170: (Length of Sequence = 398 Nucleotides)

TTEACCICAA CITACIGAGC AATGCCGIAG CIATGGAATA GAAGCATITG TIGCACTCIT TITGIGAGCC AGGCCCTGIA
GGAGGGATTG TGGATGGCAA AACCTCAGGI TCTGCCCCAAA TCCTCCCCTT GGGGGCTGGA GGGTCTCTAG TTAATTGGCA
TTCCCGGIGCI TAAGGCCACT TTTGGGTAGA GGITTGGCAA GGATGGAGTG TCCAGACCTA TGATCCTCTA AGAACTTTAC
CTTTTTAAAAA CAGCCACCCA AATGGTGGTG GCGTGGGGAG CAGGTGGTGG TGAAGGGACT GGGGGTGTCT GGCCATKGCC
ACGTACCAGA GGAGACTCTG TGAGCCCTCT CCCTGCCTGA GGGACACTTA ACTITTATAG CACTACATAG GGTCAACG

SEQ ID NO:171: (Length of Sequence = 321 Nucleotides)

AGACAGCATC TGGCTCTGTC ACCCAGGCTG GAGTGCAGTG GCGCAATCTC GGTTCACTGC AACCTCTGCC TTCCAGGTTC

AAGTGATTCT CCTGCCTCAG CCTCCCAAAT AGCTGGGATT ACAGGCATGT GCCACCATAC CCAGGTAATT TTTGTATTTT

CAGCAGAGAC GGGGTTTCAC CATGTTGGCC AGACTGGTCT CGAACTTCTG ACCTCAAATG ATCTGCCCAT CTAGGCCTCC

AAAAGTGCTG GGATTATAGG TGTGAGCCAC TGCGCCTGGC CCTTGGGTAA ACACTTCAAA TGCAMCCAAC CATTAAAGGT

SEO ID NO:172: (Length of Sequence = 293 Nucleolides)

.....

GAAACTTATA GICTIGCCIC CCAACCITCI GAACACTCCA GIAGAAAAAT CITCICGCCI ACCITTATCA CCCCACGACC
TACTAGCATT TCITACICIC AAAAAAAAATC TITTCIGAAA AATCAAGACA GAGIGCAAAC AATCAGCATA ATTTTATTAT
GACARAACTT TTAAATTTTA TCCCCCTCIC TGAGAGKICI GCIAGGACTC CITCAGATAA GIGAAAAAAGA AAKTTTTTAA
AATTTATTCI CAAATCCGAA TTCCAATCIG TATAAAAAGG GCGATTCICC CTC

SEO ID NO:173: (Length of Sequence = 282 Nucleotides)

GCTTGGTCCC GITCCTCAGG AAAAGGATGG ACCTTCTCTT CTTCTCAGAT GGTCCCTTCC ATTCCCCTGA AACCTGCATG
AGAGCTCCTA ACATGTTCT CCAATGCAAT CAAGCCTAGA CTCCAAATGT CCTCCCAGCT CACCTCCATC TATGCATCTC
ATCTCTGGAT TTGGTGATCA GACTCTATAT TGACAGTAGG ATCTCAAACC CTGCATCCAT CCTTCCTCCA GCAAGCCCTG
CTAGCCACAT GAGGAACAAG TTTCCGTGTC TTCATGACTT CC

SEO ID NO:174: (Length of Sequence = 353 Nucleotides)

CAAGAGGIGG GAGAGGIAGG GGGCAACTAC AGCTCCCCAC CAGCCCCACC AGGGGGAAIG GACCCCTCCC TGCCTCCTGC
CCAAGIGGCT CCCCCTGIAT TATGGGGGGG ACTITGIGCA AACTCIGCCC CGAGGGGGIG GGGAGGGIGG AGGGIGAGIG
TGAAATGGCA GCGGTTGGGG CTGGCAGCTG TGCTACTGGG CACTGGGGGG CTTGIAGGGC TCCAGGAGGA GGGCCGAGAA
GGIGTTGACC TTGICTGCCC CCCGCACCTC ATGGGGTAAC AGCGGCAMTT TCACGATGIG GAAGTTCTTC ATACAGGICC
TCCAATCTGG TCCAGATACT TGGCCTGGGT TCT

SEO ID NO:175: (Length of Sequence = 394 Nucleotides)

GCCCATGCCC TIGIGIACAT AATCTCTAAT ATTTATATAT ATTGATATAG AATTCTCTCT ATAATATATG TCATAGAATC
TCTCTTGGGC CIGGCGTGGG AATGTGACAT TAAGAAAACA TGCTAAGACT GGCCAGAAAA ATGGATATTT CCCAGACCTG
GAGGATGGTG TGTGGGATGT ATAGGTGAGG TCGTGGAGAA GATAATAAAC TCATTCCCCA AGATACCCTC TTCAACACAA
GGACAAGAAG GAAGGTGTGT GGTGGGGGAG GGGACAATGG AGGGGGAGGA GTGGAAGATT TGGATTTTCA TTTAATAAAG
TCAATTGAAA AATGAAAGTG CACCCCCCT CCAAAAAAACA GGAGATTCAT TTAGCAAGAG CCGTTTCATT CACA

SEO ID NO:177: (Length of Sequence = 381 Nucleotides)

ATTGGGACGG GCCCCCCTCT GAGGGACGG ATCGATAAGC TTGATATCGA ATTCCTTGAT NITITCTAGT GFTATGGTTT
TCTCCCACTC CAATAACTWT TCATACCTKT GGTCTKAGTT TTTCCATCTA TAAAATCATG TGCTAAATAA TTAACTATCA
TCTCTATCAT TGTCAGACTA CACAAAGCTT CCAGCCTGGG CAACAGGAAC CCTGTCTCTA AAAAAAATAC AAACATTAGC
CAGGTGTGGT GGTATGCGCC TGTATTCCCA GCTACTTGGG AGGCTGAGGT GGTAGGACTA CTTGGGCTTT AGAGGTCAAG
GCTGCAAGTG AGCTGTGATT GCGCCACTGC ACTCCAGCCT GGGCAACAGG GCAAGACCCT G

<u>SEO ID NO:178:</u> (Length of Sequence = 443 Nucleotides)

GATITIATIC AAACACAGGC AAGAACAATG ACCTICAGAG CIGGGIAAAA ATAATAAGIT AAAAGCATGG TIAGAATITI
AGACCAATCAG ATAAAAAGIT TGAAGGAAGI GATITCCCCI TCCTCTCCIA ATTGATTAAT TCAACACAGC ATAAAAATAA
TTIGIATCIA TAAAATATCC TTGITCCCAC ACAAATGAAC TGGAGGIGGC CCIAGGATIT CCTTGACTAT GCACAATGCA
CACAATCIAC ATGICCCTCC TCCCCAACIT TTAAGGCAAA AATGGICCTG CATCITCAGG CAGAGGGIGG GCTCATGCCA
GCAGICAGCT GIGGICAAGG ACACTGGGGG TGCGTTTYCT CCACCGAAAG ATGCCTGCTT TGGGTCCACT TTGGGCGCGG
GATCCCATTT TATTTTCTAG CCTGTGCCTC ACCACAGGGA AAA

SEO ID NO:179: (Length of Sequence = 325 Nucleotides)

TGGGGGACCA GCATTGCTCC CAGCTGAGGG CGCCGTCTTC CTCACCACGT ACCGGGTCAT CTTCACGGGG ATGCCCACGG
ACCCCCTGGT TGGGGAGCAG GTGGTGGTCC GCTCCTTCCC GGTGGCTGCG CTGACCAAGG AGAAGCGCAT CAMCKTCCAG
ACCCCTGTGG ACCAGCTCTT GCAGGACGGG CTCCAGCTGC GCTCCTGCAC ATTCCAGCTG CTGAAAATGG CCTTTGACGA
GGAGGTGGGG TCTTACAGCG CCGAGCTCTT TCCGTAAGCA GCTGCATAAG CTGCGGVTAC CCGCCGGACA ATCATGGCCA
ACTTT

SEO ID NO:180: (Length of Sequence = 213 Nucleotides)

GAGCATGCCC CCGGAGTCCC CAAGATCCTG GTGGGGAACC GCCTGCACCT GGCGTTCAAG CGGCAGGTGC CCACGGAGCA GGCCCAGGCC TACGCCGAGC GCCTGGNCGT GACCTTTTTT TAGGTCAGCC CTCTTTGCAA TITCAACATC ACAGAGTCGT TCACGGAGCT GGCCAGGTTC GTNCTGCTGC GGCATGGGAT GGACCGGCTC TTG

SEO ID NO:181: (Length of Sequence = 219 Nucleotides)

AGCITTATCA CATTATACAC AAACATAGAA AACAGTGTT CAGAAGAGAA GCAAAGGCCA TTGGCTTCAA ATATTTATGC AACAATGAAA ATGTTCTCAG CCCTTAAATG AGCACTTGTG ACTTGTCCAA CAGTGAGATA ACTAGTCAAT GGAAGAGTTC AACACTAGAG CATGTATCTC AGTCTGTTCT CATATTGCTA TAAAGGGCTS CCTCAGACT

SEO ID NO:182: (Length of Sequence = 451 Nucleotides)

GICTIACICI GITACCCAGG CIGGAATGCA GIGGIGIGAT CATAGCICAT IGCAACCICI GCCCICTAGG CICAAGIGAT CCICCCACCI CAGCCICCG AGIAGCIGGG ACTACACGIA CATGCCACCA IGCCCAGCIA ATTITIGITAT TITIGGIAGA GACGGGGITT IGCCATGITG ACTAGGCIGG TCITGAACTC GIGAGCICAA GIGATCIGCC IGCCTCGGCC ICCCAAAAGIG CIGGGAATAC AAGCGIGAGI CATGGIGCCI GGCCTAGITI GCICTTATIT TITITCCATC TITGCAGITI CIAGGCCACT GGGAACAGGC IGCAGAGCIC AGAGICCACA GCIGIGAGGC TCCATGITGC ACCATCAAAA AATAAGGIGA CGAGAGTCCT GGGTTTCCCA GIGICACGGC AAGAGGGGIT ACTGCTCACG GGIACACACA G

SEO ID NO:183: (Length of Sequence = 444 Nucleotides)

CCAAGITGAC CCGCCGAACC ACCGAC.GGA AGAGTGAGIT CCTGAAAAACT CTGAAGGATG ACCGGAATGG AGACTTCTCA
GAGAATAGAG ACTGTGACAA GCTGGAAGAT TTGGAGGACA ACAGCACACC TGAACCAAAG GAAAATGGGG AGGAAGGCTG
TCATCAAAAAT GGTCTTGCCC TCCCTGTAGT GGAAGAAGGG GAGGTTCTCT CACACTCTCT AGAAGCAGAG CACAGGTTAT
TGAAAAGCTAT GGGTTGGCAG GAATATCCTG AAAATGATGA GAATTGCCTT CCCCTCACAG AGGATGAGCT CAAAGAGTTC
CACATGAAGA CAGAGCAGCT GAGAAGAAAT GGCTTTGGGA AGAATGGCTT CTTGCAGAGC CGCAGTTCCA GTCTGTTCTC
CCCTTGGAGA GCACTTGCAA GCAGAGTTTG AGGCTCAGCA CCGA

SEO ID NO:184: (Length of Sequence = 399 Nucleotides)

GGCAGAAAGA GGAAGGAGAC AGTGCCAGGA GGAAGAAGGA AGGAGTCCCT TAGCTCTCT CATTGTCCCC TTTACTTCCT
GCTATCTTCT TCTCCTCTC TTCTCTCTCT TGCCTNTATG CCTGTATTTC TGGCAATATG ACAGGCCTGC CTACCCAAGA
TCAGAACTCC AAAACCACTC CCACCCCTGA AGGTCGGGAG GGTCTTAGCA GCCCTGGGTG GCTGCCTGTG CTCAGGTCCT
CAGCTCCATG GGAAATAAAA ATGGCACCCT GAATCTCTAG GATTTTGTCA CTTTGGAGTC ACAGCAAAGT TCTCTTCCTC
TTGTCCCCCC GTTTGCTGCT CCTTGGGTTA TAGGACATGG TAAATATTTA TTACTTTCAG GGAACCAGTA TTTTATTAG

SEO ID NO:185: (Length of Sequence = 263 Mucleotides)

 TCAGCAGAAG GCTCTTGGGC GTGTGAGGGA GGGTCTTGGA GAACTAAGCG AAGGAGGCAA ACGCCAGGGC CCCTTGCAGGCACC ATGTGCACCA CTT

SEO ID NO:186: (Length of Sequence = 343 Nucleotides)

GITCCAATAG CIGGITITAT TCICAGCACA AAAGGGCCCI GIGIAAAAAC CAGAAGGAIT TIGIAAAATA TCAAAATGAA
TATITGGCCI GGAGGITGGA AAGIGAAGCA AGGCIGGACA TAGAAAAAAA CIGATCAGIA GITATICAGG ATAITATITA
GGATAAATGA AATAGGAACT TAGGGGCATC TCITACITIT CIACAGGITC TIATCIGGGI CAATGAAGAA AITGIGITTA
TCITGCIGCC CITGCATCAG GITITITGCA CIAATGGAAA AAAGCCGGCC GAAAAACAAA ACCCAATCCI TICAGICCIA
GCITITACAT CITGCCCTIG CAA

SEO ID NO:187: (Length of Sequence = 229 Nucleotides)

GGTGCGGCTC CACCCCTTCC ACGTCATCCG CATCAACAAG ATGTTGTCCT GTGCTGGGGC TGACAGGCIN CAAACAGGCA TGCGAGGTGC CTTTGGAAAG CCCCAGGGCA CTGTGGCCAG GGTTCACATT GGCCAAGTTA TCATGTCCAT CCGCACCAAG CTGCAGAACA AGGAGCATGT GATTGAGGCC CTGCGCAGGG CCAAGTTCAA GTTTCCTGGC CGCCAGAAG

SEO ID NO:188: (Length of Sequence = 284 Nucleotides)

CCAGCAACTC AAATTCACCA CCTCGGACTC CTGCGACCGC ATCAAAGACG AATTTCAGCT ACTGCAAGCT CAGTACCACA
GCCTCAAGCT CGAWTGTNAC AAGTTGGCCA GTGAGAAGTC AGAGATGCAG CKTCACTATK TGATGTACTA CGAGAKGTCC
TACGGCTTGA CCATCGAGAT GCACAAACAG GCTGAGACCG TCAAAAGGCT GACGGGATTT GTGCCCAGGT CCTGCCCTAC
CTTTCCCCAAG GAGCACCAGC AGCAGGTTTT TGGGGGCCAT TGAG

SEO ID NO:189: (Length of Sequence = 215 Nucleotides)

GGAAGGATGA GAAACAGATT TCTGCTCACT TCATGGGCTG RCCTRGRATT GACGATGGTR CAAACCCAAG ATTATCCTCA
TGTAATTTAT GAAGATTATG GAACTGCAGC GCATGACATC GGGGACACCA CGAACAGAAG TAATGCAATC CCTTCCACAG
ACGTCACTGA TACAACCGGT CGGGCACATC TCKCGGCCTA TGCTGCCGGT GGTGC

SEO ID NO:190: (Length of Sequence = 153 Nucleotides)

TITICATATIGG AAAGAGCTAG TACAATCACA TATTIGAAAG GAGAAACAAT AGGTACTGAA CCGGAGGGAA AGGGCGAGGG
TGAGTGTGCC AGCACCGGCC TGGTGAATCC ACGATTCGGT TTCCCCATCCA AGGGTAAGTT TCCCAAAATA CCG

SEO ID NO:191: (Length of Sequence = 316 Nucleotides)

GIATTIATAC ATTIATITAT ATATGIATAT TIACTICAGA NGAAACGAAC ATTICGGGGA CAGGAAGCAA GCAGGCCCGG
GGCIGCTICC CICACIGCCC ACCICAGAGT CAGAGTITGGC ACATGACAAA TACCAAGCTC AGGGIGAAGA ACTGGGAGTI
AACTGGGAAG TAGGGRGCGC TCTATGCACA CGCAGGCTTC TAAGGGTGCA CGGTATGGGC AGKKGGTTTG CACTGGGAGG
CCCTATGTAC AGCTTGAAAG CTAGGGGTGA GATTAGCCCA GTGACTACAG GAACATACGT CAAAGTTGAG AGAAGA

SEO ID NO:192: (Length of Sequence = 360 Nucleotides)

GIGGITTITG GITATATGCA GCTITTGACT AGCATGIATT GIGICITTIT CICCICIATG AATAATTITA TATTICATGC
TACTICITGA AAGITTACTC TITGATGCTC TAAGAGAACA GCCAGATGGT TIATATGAAT AANCIITATC TGCAGGATGG
TGGATTGGTA AATNAGGAGA ATGITGITTG AGATATCAAG ATTITATGICT GGGAACTAAA ATATATAATG CCAAATGIGT
TITTIGICAAT TACTAGAGAA TICTGIGCAA ACATATCATC TCTTCACATG CTGCACACTT TGCTTTTTTGT TAAACAGCAG
GTAGTAGACA GACCAATACC AGTITCGCGT TAAGG

SEO ID NO:193: (Length of Sequence = 397 Nucleotides)

GAAAAGACCA AGGAGATGGT GAAGACAGCA GAAGCCCAGA AGCAGCAACT GAAGGAGGAG CAGGGGAGGT CAGCAAGGAA
CGGGAGAGTG GGGATGGAGA GGCTGAGGGA GACCAGAGNA CTGGAGGGTA CTATTTAGAA GAGGACACCC TCTCTGAAGG
TTCAGGTGTA GCGTCCCTGG AGGTTGACTG TGCCAAAGAG GGCAATCCTC ACTCTTCTGA GATGGAAGAG GTAGCCCCAC
AGCCACCTCA GCCAGAGGAG ATGGAGCCTG AGGGGCAGCC CAGTCCAGAC GGCTGTCTAT GCCCCTKTTC TCTTGGCCTG
GGTTGGCGTG GGGCATGCGT CTAGCTTTCA CTCTGGTTCA GGTCCAACAG GGTCCGTTCT GTGCCTTTGG TGCCCCC

SEO ID NO:194: (Length of Sequence = 225 Nucleotides)

GATTATTGGC TTTGCTTTCA TAACATGTAT TTTTAAGTAT TTACTCTCTT AATGGCCCTC GTGTCTATTT TATACATCAT
ATCTCTTAAT TCTCTAGATG GAACACTGAA GGACAGGAAT TAAGTAAGTG ACTGGCCATG CAAGGGTTGG AAATTTTACT
GTATCCCTTC CTCRGTAGAA GTTATGTTAA ACATTCAAGC AACCACATAT CTAACAGAGG AGTTT

SEQ ID NO:195: (Length of Sequence = 294 Nucleotides)

ATTACTAGAT ATTIGTATGT TAAATTATGT GGGTTTCAA ATTIGTGGAG AATAAGTAAT AGTGACATTA GTTTAAGGAC
AGTGTTTCAT CAGGGCATTA TTTTAATGAA TCTTATATTT AAATGTCTGT TTCAGGAATT CATGTGAATC TTTCTTTTTA
TAGAGGACCC ACAGGCATGA NITATTTACT CCTCCGGTGA TAGGTTCTCA CCCTGATGAA AGCGGAAGCA AATTCCAGGT
TAGAACATTA TNCTAGTTAT GTAGGGGGGT ATAAAGTGTG TAAGTTTAAT ATTT

SEO ID NO:196: (Length of Sequence = 233 Nucleotides)

TTATTTTCT CTAAATTTTA AAATAGAAGA CTTTAATGGA AAACATTTAG TACCATCATG TCAMCCTGAA TGCCAGCAAT ACCTCGACTT TTACACACGC AGGAAGCCTA GTAAAAGCCC CGTCAGTAGT ACACATTTCT CTATGGTCCT TCAACAGTTT TTCATATACA AAATTTTCTG CTATTTTTGC TTTTGCAAAC AGCAATAACT TTTGGGTTTC CCATATGACC ACC

SEO ID NO:197: (Length of Sequence = 230 Nucleotides)

AAGATATCTA CCTGGAGTAG CTGTGCAGCC CCGCCCTCTG CTTCCCCCAG CCCTCAGGCC AGTGCCAGGA CAGCTGGCTG CTGACAGGAT GTGGCACTGC TTGAGGAGGG GCACCTGCCA CCGCCAGAGG ACAAGGAAGT GGGGGCCGCT GGCCAGGGTA GGGAAGGKTG GGGCAATGGG GAGAGGCAAA TGCAGTTTAT TGTAATATAT GGAATTAGAT TCATCTATGG

SEO ID NO:198: (Length of Sequence = 118 Nucleotides)

TTCTCCTGGG GAAAGGCTG TTGCTGAAGT GGCCGGTTT TTTAAGCATC GACATTTGCA TCCAAAGGTT CAAGCAGCCG CCTCAGGTTC CARAGGCTTC CACCTGATGG CTGCACTT

SEO ID NO:199: (Length of Sequence = 268 Nucleotides)

TAAATGATGG AGITAAATGA TGTTGTCAGT GCCTATTTAA AAAACTACTC TTCCCCTTCT CTATGAGTTC TACTTTGGTA
AATATTAATA TTTAACCAGT TAGTAAAACT AACACCACTA TTTCAATTCT CTTTTGTGCA TAGTAAGTAA ATTTTGCTTT
ACTTACTTTA TAAAAAAAATA CTTTACATTT TATAAAGCAG GTTTTAGAAA AACGGTTTAC AAGAAAGTTT GCCTCCATTT
CACTGCCAAT TTAAGCACAG GGGAAAAT

SEO ID NO:200: (Length of Sequence = 422 Nucleotides)

CCAGTGAGTT TGTGAAAAGC AACAGGGGTA NGACAGGTTC AAGGAAGGAC ACAGACAGTG CCCTGTTTTA CGTTCCAAAT TTCTTCTTTT TAATGGGTGG TGGGAGCTGA GCAATGATGT CATTTGGAAG GGGCAATGAC TTGTCAATNA TGCAGAACAT GTAGGCATCA TGGAGAAGGA TGTGCATCGG TCTCTTGGGA TGAAAACTGA TGTGTGTGAT AAGAACTATCC CTTTXTAGCC

· . .

AAAGGIGGIG AAAGCCCIGC TICIGGACAG TCCGGCTCCA ATCTGIATAC TGITTGICIG GGATGCIGIA CTCAAATACC TGCIGGICCG AATGAGCGAT GACAAGGIIG TTTGGIATIG GGGGCAATAG CCATAGCAGI CACTIGGGAA ATTGIAAGCA GGCACCGIGC AGIGAAGIIT TA

SEO ID NO:201: (Length of Sequence = 273 Nucleotides)

ACTOCACGOT GATGAACCOG ACGTCCATTT CTCCAAGAAA TTCCTGAACG TCTTCATGAG TGGCCGCTCC CGCTCCTCCA GTGCTGAGTC CTTCGGGCTG TTCTCCTGCA TCATCAACGG GGAGGAGCAG GAGCAGACCC ACCGGGCCAT ATTCAGGTTT GTGCCTCGAC ACGAAGACGA ACTTTGAGCT GGAAGTGGAT GACCCTCTGC TAGTGGAGTC CAGGCCCCCA GACTACTTGT TACGAGGGCT ACAACATGTG CACTGGGTGC CCG

<u>SEO ID NO:202:</u> (Length of Sequence = 436 Nucleotides)

GGACTICCAAC CCCCCAGGAG GCCGAATGCT GAGCTTGGCA ATGGTGGCCT GGATGGAGCT GATGGGCACA TCCCCACCGA GGACCAGGTC CTGGGGAGTCC TGAGGAAGGT GGTTCTTCTG GCTGATGCTT GCACTGGCCA AGGGTTTGCA TGGAGGAGGC ACACCATGGC GCTGCAGGAC CTGCTCCACG TGTCTCACCA CTGCCTCATA GCAGAACCTG AGGTGCAGGT TCTCCTGCAG CATGTGCTTT CTCTGCTGCC GCATGCGCCG CACCAGCTGA GGCAGCTCAG GGATTCCKTT CCCAGCCTCC ACCTCCTGCA CAGCTGCATA GAGCAGTGCA AAGGCTCCCG TGCGGCCCAC ACCAGAGCTG CAGTGCACAA TGATGGGCGT TTGCAGGGGC CGTGATGCAA GGTAATTTGC GTGCACCTCC TGGGTT

SEO ID NO:203: (Length of Sequence = 336 Nucleotides)

CIGCATGINT TGGGGACACT TACGCCAAGG CGCCGCGITC TCATTAGGAG CIGGGACCAG AAGIGAATAA GCCAGGITCC TGTCTCAGGG AGCTCCATAG CAGGACTCAG AACCACACAC GGCCCTCTAG GCATTIKTGA AGCTCTGTGC TTCATTITTT TTGCTTTGCC TCTAGTTTTG CCTTTGCAGT ACCAATGCAG CCAGCCCATG TKTCCCCTCT ATGTGGAATG TTAACGATAT TCCCACTGTT TCTGGTGTCC TTTCTGTAAT CAGAGCTGCC GTGACCATTC CAGTTCAGGC ATCCTGGTGG CCTGGCTTTC TCTGGGGCCAT AGAGCT

SEQ ID NO:204: (Length of Sequence = 393 Mucleotides)

GERATCAGAT GCTCAGGTGT CCAAGCAGGG ATAAGGACAG GCAAAATAAA TAACCCCCCA ACCCCCATCG TCACTCTGCT GCAACACGAC ACAAAGGTTT AAAGATCTGG GCCCAAAGAC TCTGGGTCCC TTCAAGCAAG CTCAGGTGGA AGGAGGTTTC CCCACCCCCC ACCAGGCCTG TTTGCCCCAG GTTGCCCTAG GATGGAGGCA GTTCAGACCC TGGGTCACTG AAGCTGATAG GAAGAACINC GATATCAATG GCCTAAGCCT GCTGINTGCC CAAGGGAGCC AAGGGCAAGA GCCAAAAGGC CAATTTAAAG GACGTGGACC TGGGGGGCCA GAGGAGGCAC CACAGCCGAG GGGAGCCACG CCCTGGGCCG GCAGGGCACA TGG

SEO ID NO:205: (Length of Sequence = 390 Nucleotides)

GAGGAAGAGG ATGACCTGAG TGAGCTGCCA COGCTGGAGG ACATGGGACA ACCCCCGGG GAGGAGGCTG AGCAGCCTGG
GGCCCTGGCC CGAGAGTTCC TTGCTGCCAT GGAGCCCGAG CCCCCCCAG CCCCCGCCCC AGAAGAGTGG CTGGACATTC
TGGGGAAACGG GCTGTTGAGG AAGAAGACGC TGGTCCCAGG GCCGCCAGGT TCGAGCCGCC CGGTCAAGGG CCAGGTGGTC
ACCGTACATC TNCAGACGTC GCTGGAGAAT GGCACACGGG TGCAGGAGGA GCCGGAGCTG GTGTCACTC TGGGTGACTG
TNACGTCATC CAGGCCCTGG TTCTCAGTGT CCCACTCATG GACGTNGGGG AGACGGCCAT GGTCACTTCT

SEO ID NO:206: (Length of Sequence = 172 Nucleotides)

٠.

CTITIACTIGIG GGTGTGGGTG TCACTGTCAC TGCCACAGCC ACTNGGAGGG ACACACAGCT TTAACCCCTR TTTGCTTAGG NGAAGGGTGG GGGCATTCAG GGTTATAAAA CTAACTATAT ACACAGAAGG TCCTAGGKAG AAAGCCACCC TGAGCACACA TGTCTAGGCA CA

SEQ ID NO:207: (Length of Sequence = 215 Nucleotides)

AAGGCAATTA GAAGATTTAT TGAATATTGG TTAAAAGTAG ATTGACAATG ACATTAAAGA ATAAAGTGTA ATTTATTTGG
TGCTACTTTG TGAATGCTTC CAAGTACAAA TCATCTCACA ATACCATATA CAACATACTT TCAATCACAA CTCAAATATA
AAATAACCTA CAAAATCACA TTGCTATAAT CAATATACAA TAATTGTATT TTTAA

SEO ID NO:208: (Length of Sequence = 444 Nucleotides)

GGAGTICTCT TGTCCACGGA GAGCAGTGTT GCAGTGTATG GAATGCTAAA TCTTACCCCA AAGGGCAAGC AGGCTCCAGG
TGGCCATGAG CTGAGTTGTG ACTTCTGGGA ACTAATTGGG TTGGCCCCTG CTGGAGGAGC TGACAACCTG ATCAATGAGG
AGTCTGACGT TGATGTCCAG CTCAACAACA GACACATGAT GATCCGAGGA GAAAACATGT CCAAAATCCT AAAAGCACGA
TCCATGGTCA CCAGGTGCTT TAGAGATCAC TTCTTTGATA GGGGGTACTA TGAAGTTACT CCTCCAACAT TAGTGCAAAC
ACAAGTAGAA GGTGGGTGCC ACACTCTTCA AGCTTTGACT ATTTTGGGGG AAGAGGCATT TTGACTCAAT CCTCCCAGTT
GTACTTGAGA CCTTCCTCCC AGCCTGGGAG ATGTTTTTTG TATT

SEQ ID NO:209: (Length of Sequence = 338 Nucleotides)

GCAGATCACT TGAGGTCAGG AGTTCGAGAT CAGCCTATAT ATGCAAGTAC ACACACAGGC ACTCGCACGC ATGCATGCTC
ATGCAACACA CATGTACACT CTACATGTAC AGCTCACATA TGCATCCATA CACATGTGCA TGCTCACCCA TACACCAGGC
ACACACAAGT ACTCATACGC ATACATGGCC ACACACAAAG TACACCACG TACACCATAT GCATATGTAT GCACTCATAC
ACTCATACAT ATGTGCCCCC TCAGAGAAGT ACACAAGTGC ATGCGCATCA CACATGCATA CGTGCTCATG CATACACACG
GGACATTTCA TACACACG

SEO ID NO:210: (Length of Sequence = 371 Nucleotides)

GAGGAAGTAG AGCCINAGGA GGCTGAAGAA GGCATCTCTG AGCAACCCTG CCCAGCTTGA CACAGAGGTG GTGGAAGACT
CCTTGAGGCA AGCGTAAAAG TCAGCATGCT GCAAGGGGAC TGTAGATTTA ATGATGCGTT TTCAAGGGTA CACACCAAAA
CAATATGTCA ACTTCCCTTT GGCCTGCAGT TTGTACCAAA TCCTTAATTT TTCCTGAATG AGCAAGCTTC TCTTAAAAGA
TGCTCTCTAG TCATTTTGGG TCTCATGGCA GTAAGCCTCA TGTTATACTA AGGGGGAGTC TTCCAGGTGT GACAATCAGG
TTATTGGAAA AACAAAACGT GGTTTTGGGA TCTGTTTGGG AGACTGGGGA T

SEO ID NO:211: (Length of Sequence = 295 Nucleotides)

CCTCCCAACG TGTTGACATT ACAGGCGTGA GCACACGCAC CCAGCCCATC TAGCATAATG TTTTGCATAG TTGTCAGCAG ATAAATATTG AATGACAAAA CTCAGATGGA GGAAAAAGAA CAAAATAACC TAGTTCTCAG AAAGATTTAA TGAGCAAATG GGAAAATGTC AAAAAGATTT ACAGACAGGG GCATCTTAGA GTCACTGGAA TCACACAGGC CTTCCCTCAG CTTGAGGGGC TGCCTGGAGG TGGGGGTGGG GGTACACCTC CTCAGTGGGG AGAGACTTGC CAAAT

SEO ID NO:212: (Length of Sequence = 370 Nucleotides)

TGGCCGATAT GAGGGGGGTG GGACTGGGCC CCGCGCTGCC CCCCACTGCA TTCTCGAGGA GGGGGGGATC CGCGCATACT TCACGCTCGG TGCTGAGTGT CCCGGCTGGG ATWTACCAT CGAGTCGGG TATGGGGAGG CGCCCCCGCC ACGGAGAGCCC TGGAAGCACT CCCCACTCCT GAGGCCTCGG GGGGGAGCCT GGAAATCGAT TTTCAGGTTG TACAGTCGAG

CAGTITIGGT GGAAGAGGGG GGCCCTAGAA ACCCTGTAGC GCAATGGGGT TGGGCGCCCC AAAGGTTAAG TTTGAACCCG AAGAGCAAAG GAAGAGGCGA TCATCATAAG TGGAGGATTA GGATTAGGAT

SEO ID NO:213: (Length of Sequence = 302 Nucleotides)

ATCTGTGGAA TAATCTGCGG GCTAACACGG ATAACTCAGT ATAAGAACCA CCCAGTTGAT GTCTATTGTG GCTTTTTAAT
AGGAGGAGGA ATTGCACTGT ACTTGGGCTT GTATGCTGTG GGGAATTTCC TGCCCANTGA TGAGAGTATG TTTCAGCACA
GAGACGCCCT CAGGTCTCTT GACAGACCTC AATCAAGATC CCAACCGACT TTTTATCTGC TAAAAAATGGG TAGCAGCAGT
GTATGGGAAT TTTCTCATAC AGAAGGGCAT CCCTCAAACC GGAAACCACA GAGATGCTAG GT

SEO ID NO:214: (Length of Sequence = 354 Nucleotides)

ATGGATGAGT GGGCACCCCG CACAGGGCTG CAGGGTGGAA AACGCTCGAC GGCCAGGTGG TGACTTGGGG GCAGAGAGCG CAGTGTNGTA GGGGAGGAGA GGTGGTGTCC CTGCTGCCTG GGAGCCAGCC TGCCTGTNCT GTGGGCAGAG CAAGGCACTT TCTGCTGCCG GTGCTTCCAG GGCCTAAGCA GCCGCTGCAC ACTCACCAGC GCAAGGCTCC TCTGCAGGGA ACGAGGGCTG CTACCCATTT CACAGATGAG GGCAAGCAAG GACTTGCCCA GGGTTGCCCA NAGCAAGTGC GTAACAGGCC CTGAGAAGAG NGCCAGTGAG CTCATCCTGA GTTAATTATG GGCT

SEO ID NO:215: (Length of Sequence = 260 Nucleotides)

TGGTTCAAAG TCTAGGCCCT CTTNAGAGCT GGCTGATTCA GCTTGCCAAC AGTGACATCA GGGTGAGGCT TCCTCTGTCC

ACAGCATTAG CTGCGAATAT CCTCATGGTC ACAAGATGGC TGCCAGTGGC CGTCAGGGTG TGTGCTTCCT TGTTCACATC

CAGTGGAAGA GTGACAGCCT GCTCCCCTTA GCTCTCTGAC ACCANTGTGA AGGTGCCANG AACTTACTAG CAGGNCTTTC

CTCATGACCC ATTCAACAGG

SEO ID NO:216: (Length of Sequence = 232 Nucleotides)

SEQ ID NO:218: (Length of Sequence = 219 Nucleotides)

CTGCAACCAT CCATACCTTT INCCCGTGGC TGCTATGGAG TCCCCCAAAC TCCCCAGTGG GGCTTATGAG GGTGGGGCAC TTATTANGIN GTCTGGGAAG CTCATGCTGC TCCAGAAGAT GCTGCGAAGC TGAAAGGAC AAGGACACCG AGTGCTCAAT NTTCTCGCAG ATGACCAANA TGTTAGCCTT GCTTGAGGGC TTTCTTAGNC TATGAGGCT

SEO ID NO:219: (Length of Sequence = 390 Nucleotides)

GATAGGTAGC AGAGACCAAG GCGCAGGGTG CTTCAGATGA GCAAGAGAAC CCAGTCGAAC CAGATACCCC AGGTGGGCCG
GAGGGACCCC AGACCTTCAG AGGGCTGCCC TGGTGTTCTC CACAGTGCAG TCCCTCTGTA TTCCCAGAGT GGGATCGGGG
CTTTCAGCCC ACCCTGATGC CTGCCCTCCA GGATGGCTGG TTTAGTCTGG GTCCATGTCC CAGACCCCTC TATTCTGCTC
CAGGACAGCA GGACTTCAGG TCTTTCCTGG GGGTGGATAT AGGAGAAAAT TTCTGCCTGG CACACACCTG GGCTCCAACC
ACTTGCCAAG TGATTCACTC TTAGGCCCAG GGGGAACACA ATGACTATCA TTACTGATGC AGACCTGGCT

SEO ID NO:220: (Length of Sequence = 382 Nucleotides)

TITITGITIT GITTIAATAT TITIGATATT CICTITGCAT TGAAATGGIA TAAATGAATC CATTIAAAAA GIGGITAAGG
ATITGITIAG CIGGIGIGAT AATAATTITI AAAGITGCAC ATIGCCCAAG GCTTTTTTIG TGIGITITTA TIGITGITIG
TACATTIGAA AAATATICIT TGAATAACCT TGCAGTACTA TATITCAATT TCTTTATAAAA TTTAAGIGCA TITIAACTCA
TAATTGTACA CIATAATATA AGCCTAAGIT TITATTCATA AGTTTTATIG ANGITCTGAT CGGICCCCTT CAGAAATCTT
TTTATATTAT CCTTCAAGIT ACTTTCTTAT TTATATTGTA TGIGCATTTT ATCCATTAAT GT

SEO ID NO:221: (Length of Sequence = 314 Nucleotides)

GACTITIGGIT TATITAAAAA ACAAGCCAAA AAAAAAAAA AAAAACCCCA ACTITATATA CAAAGTCAAA CTGAAACCAC
GGWITATGGA AAGAGGCAAG AWITATGGGT AACAGGGGAG AAGCCTGGGC CAGAGCCAAT ACCACATTCT GAACACAGGA
GCCACGGGAA AGAGGTGCTG GTTTCTTCTG GCAAGACCGG GGTGACTGGA ACGCAGTGGT CCTACTGGCA AACCCAGCCC
AACACTGAGC TCTTTCTAGC ATGGACTCCA TTCCCGTGAT TGGCCAAGGG AGACCCTTCC CCCAGGAGGC CTGT

SEQ ID NO:222: (Length of Sequence = 342 Nucleotides)

TTCCTTCTCT GCGGCGCAC GTCGCNAGCA GCCTGCTTCG CCCCGTCGTC AACTITGAGC TGGAGGAGAA GCAACTTTGG
CAGTGGCCGC GGGGTGGGAA TCCCGCTTCT CCTCGGCAGC AGTAGGCTCG CAAGTCGCTG GGGTTAGGTG GGGCAAGAGT
TTCGCCGGCG CATCAGCGCT TGCTTCGGAC TGTTTGCAAC GTGTTTCCAG CGAGCTGGGA GCGGGGTTG TGACTGCGAG
TCGTCTGGGG GAGGGGGACT TGTTTTTCTT TTCCTCTAGA GACCTCGGCT TTCAACTGGA TCAAACGTTG TCGAAAGGAT
GTAAATAGGC AAGAGCAAAC TG

SEO ID NO:223: (Length of Sequence = 376 Nucleotides)

GIGATGGCTG CCTTGAGGGG GACCATCATG TCGGAGAGGGC ATTGGTGCAG GTCTCACCCC ACAGCCCATG CCCAGCCTCC
TGCAGACTCA GGTCATCCAG CTGGTCGATG GCTCTTTGCA TACCTGGTGC CTTCTCCTCT CGGGCTTGGC AGGCTTCTCT
GGGGGCTTCT CAGATGACTC TTTTGCCTTC TTCTCTGTCT TGGCTAACTC CTTGGCCAGC TCTGAACGTG CCTCCTTGGC
TCCCTCTTCT ACCACCTCCT CCCGTTTGGC CAACTTGCTC ACGGCCGTCT TGGTAGTGGC TTTGAGGCTC TCCTTGCTAT
CAGCCCGCTG TTTGATTTTG CTGGGCTTGA GGTTGGTAAG GCACAGCCCC AAGAAG

SEO ID NO:224: (Length of Sequence = 445 Nucleotides)

GTTGATAGAC ATTGGCATTG GGGTTGCTTC CACCITITGG CTGTCATGAA TAATATTGCT ATGAACACTA ATGTACAATT
CTTTGCCTGA ACGTAAATGT TITCATTTCT CTTGGGTATT TATCTAGAAA TGAAATTGCT GTATGTTAAC CCTTTGTTTA
ACCTCTTGAG GAACTGGCAG ACTTTTCCAA AGCAGCTGCA CCATTTTAAA TTCTAACCAG CAGTGTTGA GGGTTCCAAT
TTCTCTATAT CCTTGGTAAC ACTTGTTATC TGCCCTTTTG GTTAGAGACA TCCTAGTGAG TGTGAAGTGG CATCTCACTG
TGGTTTTGAT GTGCATTTCC CTGATAGCTA ATTGTGGGA TCCCTTTTGC TTTTAGTGGA ATGAAATATC TGGTAGTCTC
GTATGCCAAA CTAAAGCTAA AATTAAAATG ACTCTGCATG ATGGA

SEO ID NO:225: (Length of Sequence = 403 Nucleotides)

TECTOTOGG ACAGITITOCO GEGCAGOTOC TEGCCAGOTT COAGCCCAGA GIOCICAAGI COAGGCACO TIEGGCCCAG
CGCAGGCAGA ATCCGAGGIG GIOCIGGCIC TACCCIGGGC CICCTACTOC COAGCACCCC TEGAGGAGGC AGGGGCICCC
CGCCGCCGAG GCIGCCTGCC CTAGGCCCAC CTCTGCATGC TGCTCATGGG GCCACCCTGC CTCCTGGGCC CTCACTCTGC
CTAGGGGAGC TGGGCCAGGC ACTAGCCTTT GCCCAGGGAG GTGGGCCTCA GGCTGCCCAG GTGCCTGCAC CCCAGCCGGG
CTTCTCTGGG GCCTCCCCGT CGTCAAGCCT ATATCCTGTC TGTCCCCACC CCAGCTGTCC CTTGCCAGGG GACTGGCATA
AAA

SEO ID NO:226: (Length of Sequence = 440 Nucleotides)

GIGCCTTAAG GAGAGAGATT GIGITCITCC TCTCTCAGG GIGATAACTC AGGAAGCCTC TGGGITGGGA AGACCATCAG
TTCTTTTGTC TTAGGITTCT TTTCCTGTCC CTCTTCCATC CCCAAGATGT GACCCCATAA AAATTTTTCC TGAGTTGGCC
AGGCATGGTG GCTCACGCCT GIAATCCCAA CACTTTGGGA GGCTGAGGCG GGCGGATCAC GAGGTCAGGA GTTCGAGGACC
AGGCTGACCA ACATGGTGAA AACCCCATCT CTACTAAGGA TACAAAAATT AGCCGGGTGT GGTGGCACAC ACCAGTAAGT
CCCAGCTGCT CAGGAGGCTG AGGCAGGAGA TTTGCTTGAA CCTGGGAGGC AGAGGTTGCA AGTTAGGCCG GGATTGCGCC
GTTTGTACTC CAGCCTGGGC AAGCAGAGCA AGACCATCTA

SEO ID NO:227: (Length of Sequence = 426 Nucleotides)

GACCAAGAAG TICCGGITCG AGGAGCCCGI GGITCIGCCI GACCIGGACG ACCAGACAGN CCACCGGCAG TGGACTCAGC
AGCACCIGGA TGCCGCIGAC CIGCGCATGI YIGCCATGGC CCCCACACCG CCCCAGGGIG AGGITGACGC CGACTGCATG
GACGICAATG TCCGCGGGCC TGATGGCTTC ACCCCGCTCA TGATCGCCTC CIGCAGCGGG GGCGGCCTGG AGACGGGCCAA
CAGCGAGGAA GAGGAGGACG CGCCGGCCGI CATCTCCGAC TTCATCTACC AGGGCGCCAC TTGCCACAAC CAGACAGACC
GCACGGGCGA GACCGCTTTG CACCTGGCCG CCGITACTTA CGCTCTGATG CCGCAAGGGC TCTTGAGGCC AGCGAAGATG
CCAACATCAG GCAACATGGG CCGAAC

SEO ID NO:228: (Length of Sequence = 278 Nucleotides)

CAGGACCAGG AGAAGATCCT GGAAGATGCA GIGGATGAGT GGACGGCCTT TAACAACAAG GITAAAAAGG CCACTGAGAT
TGITTIAGAA AACCAACAGC AAAACACTGA CAAGGIACAT AAATACAGAT TGGACATITT AGGGTAAATT CACTGTATIT
CCTACTIGCT TGIAGGAAAC CGAGIAAAGT GGAAAAGCTG TCCTGATCAT ATGGCATGCA CACCAGACTG CAAAAGGNCG
TCCACACTAT TTAACAGGAC TGIGGCAAAA TAGCTTTA

SEO ID NO:229: (Length of Sequence = 425 Nucleotides)

THITTGITCC CAAGCCITTG TGACTGACTT TAAATCCTCT CACCIGCAGA ACAGAGATGG CITCAAAGIG GGGAGIGAGG
GAGTGAGCGA GGACCCTGGG CIGAGACCTG TTTTTCTTCC ATTITCTGCTG TGGCTTCCCA CAGCTCCCTG GTTCCACACC
AGGCCCTGCT CIGCCGCAGA AAATGGATTC CCAGGCCACA GAGCTGTCAG GCCTTTGACT TTGCAGAGAC CAAGCACCCC
AGAGGCTGTG CGACASGGCT AGTCCCTGGT GGGCCGGTCT GGGGCATGGG GGGCAGGGAG ACTKGGAGAT GGGGAGGGCG
TTGAGAATCC GGGGGGTCCT GGATACTTGA CAAATTGGCT CAGGTCTTAG CTYTGGYTGC CCCACTGATT GTGTTGCTTG
GCAAGGTGCA AGTYTTCGGC TGTTC

SEQ ID NO:230: (Length of Sequence = 382 Nucleotides)

TTGGAGGATG TGCTGCCCCT CCTGCAGCAG GCCGACGAGC TGCACAGGGG TGATGAGCAA GGCAAGCGGG AGGGCTTCCA
GCTGCTGCTC AACAACAAGC TGGTGTATGG AAGCCGGCAG GACTTTCTCT GGCGCCTGGC CCGAGCCTAC AGTGACATGT
GTGAGCTCAC TGAGGAGGTG AGCCAGAAGA AGTCATATGC CCTAGATGGA AAAGAAGAAG CAGAGGCTGC TCTGGAGAAG
GGGGATGAGA GTTCTGACTG TCACCTGTGG TATGCGGTGC TTTTGTGGTCA GCTGGCTGAG CATGAGAGCA TCCAGAGGCG
CATCCAGAGT KGCTTTAGCT TCAAAGGAGC ATKTTGACAA AGCCATTKCT CTTCAGCCAG GA

SEO ID NO:231: (Length of Sequence = 398 Nucleotides)

GAGGCTGGAG AATCGYTTGA ACCCAGGAGG COGAGGTTGC AGTGAGCCGA GATGGCGCCA TTGCACTCCA GCCTGGGCCA GAGCAAGGTT CCTTCTCAAA AAACTTGGAA ATCTGTTGGG AAGTAGGGG AGGCAAGGT TAAAACCTAT GCAGGTGTGT CAATTAGACT TGTTCCAACT TGAGAACCTG AATTTTGCAT GTAATTGAAA TGTTCCAGAA CAAGTCTGGC AGTTTCATAA

5,80

GGGAGTTTTT AGATGCCAAT ACATTGCAGA TAACCATATT GGTTACATTA GGGGAATGAG CATGGGATAG GTGCCTCCCA GTTGGTAGGA TAGCATGAGG AGGTTTCAAA AGTAACCSCT TTAAGGGTTA TGTCCAGTAT TTGCTAAGTA ACCAAGGT

SEO ID NO:232: (Length of Sequence = 272 Nucleotides)

GOGGCTGCAG ACTGAGTTAT TITATITCGC TATTICCAGT TIGAAGCTAC TATCATGGGC GITTAGAGTT ATACAAATGA
CACTTACAAA AAATAAAAGA CCAAGACACC CAGAGTGAGA TGCATGTTGG GGACGGGGGA GGCTGGCAGC AGGGGGGCCC
CGGCGGYTCA CCCCAGGGCT CCCGGAGGGG CGACGCCTGG CTTCATCCAC CCGGGAGGCC CAGGGAGCAC CAATCACAGC
AGGGGCTCTG GCCCAGGTGT CGGCAGCCCA GG

SEO ID NO:233: (Length of Sequence = 364 Nucleotides)

ATTITACAGT TITATITITA AATCATITAC ACATATICAT ACAAAGAAAA ATAAATITCA GGATGGAATC CIGGGGACCA
TGGTAGITTA AAAAAAAAAA TCTCTCTGAT CATTAGCTAC TAAAGACANG GCAAGAGGCT TAGCAGTCAT TICTGGGGGT
TAGTGTATCT CCCCATGCAG GGGACAACTG NGAAGAATCC AAGCTGCTCC CTCATCTTCC TICGATCTAG ATGGGGGAAG
GGGATTTTCC AATGCTCTCC CCTAGAAACA TITCAAGAAG TACAGCAAAG GCTTATGGTA ACACTGGAAC CTATTTGCTA
GAAATCTGGC AAGATTGCAC TITCTGAACC CAATTTTCCT ATAA

SEO ID NO:234: (Length of Sequence = 217 Nucleotides)

GGCCAGGAGC CAGAGGGCCC CGGGGCCACC CCTGCCGGGG AACGTGATGA CCAGAGTCCA GACAGTGTCC CAGAGAGGGCC GGGGCCCGCA GACCGGAGGC TCTGTCTGCC CTNCGTGGAC GCCTCGCCAC TCCCAGGGAG GACGGCCTGC CCGTCGCTGC AGGAGGCCCAC GCGGCTCATC CAGGAGGAAT TTGCCTTCGA TGGCTACCTG GACAATG

SEO ID NO:235: (Length of Sequence = 221 Nucleotides)

AACTITAAAG TTAGGATTIT AAAATATTIG TAACIGGCTA AATTITAAAG TOGIGACAAA TAATTACITA GGITCAGAAA TATACACACA CTTACICITT AGCCAGITIC TITCAAGGIN TTACIGICCC ATCAGATATC TAGCCATTIK CCTTIGCAAA TTACATACCT TCTTAAGAGI GIATTITTAA GATTATTACT TATGCTITTAT GATGATATAG T

SEO ID NO:236: (Length of Sequence = 221 Nucleotides)

ATAAATGGT TICTCACTCC TIAGGGACAC GATTGGAAAC AATACATCCC ATGAACACAG GTGAATGTCC CTGGTTATCC CTGAGCTGGG CAGTTTCACA CAATCANTTT TNCTCTGAGG CCAAAGTCTG TGGTTTGATC ATCTTAGCAG CTTCCAGAAC AGAAAGTAGG TTTACTTTGT CTCCAAANTC TNATTCTCGG TGCTCAAAGA AGAATGACCT G

SEO ID NO:237: (Length of Sequence = 251 Nucleotides)

GACATCITIC TAAGATICIC TGIGGGAAAA TGACTGICAA TANAATGCGG GITTCIGGGC CATTCGICIT ACTITCATIT
TTTGATTACA AATTICICIT GACGCACACA ATTATGICIG CTAATCCICI TCITCCTAGA GAGAGAAACI GIGCICCITC
AGIGITGCIG CCATAAAGGG GITTIGGGAA TCGATTGIAA AAGICCCAGG TTCTAAATTA ACTAAATGIG TACAGAAATG
AACGIGTAAG T

SEQ ID NO:238: (Length of Sequence = 327 Nucleotides)

GITCGIGGCT GICACAATAA IGCIGIGATA AIGCIGIGGI ITCCCAGCAG GGAGGIGGGA GCGGGGAGGG GGCIGCAGCC IGAIGAGAGC CAGCIGAAGG AAGAGCIGCC ICICCCITICC TAAGCCCCITI CCCAAGGICI GCCCCACCGC CCAAACCAAA GACCACICCG AACAAAGIGA GGAIGIGGAI GCICTIGCIG GGICCGCIGI ICCGCAGAGG GAAAGAAAGG GIAGCIGCAC TGACCCCACT GTCCCCATAT ACAAGGGTTK GGGGGCAAGA GCATGTGGCT ACTCCCAGCA AGGGRAAAAT GGGAGGAGCA GTAGAAA

- SEO ID NO:239: (Length of Sequence = 285 Mucleotides)
- ATTATIAGIT TATGGIGCTI TAAACCIATC AAAATAGIIG TAAGIAAATG GATITCIIGI NCICCCAATA ACAATTCICT
 GAGCIAGGAT AGAIGICITI CIGGCCATII TACAGGIGAT GACACIGACA TAGGGACIGA GIGGGIAGCI TAAGINCCAT
 GGITACCAGG AGCAGGACCN ACGITTCCIG NCICCCAGIC TCATCCIGIT TICCACIGAC CAGGITGGII GCICCCIIGG
 AAAGCAGICC CIGAGAGIIG ACTITAGAAGI TCAGGGRGAA GAGGI

SEO ID NO:240: (Length of Sequence = 349 Nucleotides)

TTTTGCCATG TTGGACAGGC TGATCTCAAA CTCCTGGCCT CAAATRATCT GCCCAGCTTG CMCTCCCAAA GYGCTGGGAT
TACAGATRTG AGCCACTGCA CCCAGCCTGA CATGCCATAG TTTCAGCATT TTCTTGGGCA ATGATCCAAG CTGAAGGCTG
GTCTGAGGGA TCTSAAGAAG CGTATGAGTT GGAAGAGAGG GACAGAAAGG AAGAAGACAT GTGAAGAGAG AAAAGGAAGG
AAGCTAGCAG AGGAATGCCC TCCAATAGAG ACTGCTGCCT GAAGCTCAGC CCCTCTGAAG ATAGGTAGGC CAGGCTGGCT
TAGCTGAGGC AGTGGGTTAG ACCAGCCCT

SEQ ID NO:241: (Length of Sequence = 233 Nucleotides)

GIGCAGCGGI CIGCCITCAT CITITAATGG COGGIGCGGI ACAGTTAGIG GACAGACGGG GGATGGGACA CAGCAGGGGI GAAACAGGGC AGTCACAGCC GGGGCCGGGG ATCIGGAAGC GGGGGCGGIC CTCCCCCTGG AAACACCGIN TCTGGAAGGA CACCCITAGG ATCCCCTGAC CICARGGIGC CACCCACACG GGCCTGGIGT TCTGGGAAGGC CCGGCTKGAG TGA

SEO ID NO:242: (Length of Sequence = 372 Nucleotides)

ATATGIACTA CATTIGGIGG AATACCCATG TACAATTCTI CAAAAATAGI AAAGAGCAAA ACAAACAAAA AATAGIAGAA GCACTGGAGA AATACACTAT GGCATAAACT AGITACGGGT GGGATGICAC ATGGACCATA TCTACACTCT GTGGCAACCT TCTTACCTGA CTCCAAAGGA TCAGATAATC AAACAGGAAA TTATGGIAGG AAATCAGAAA ATTGAAGTAT GCATTCATAT CCTAAGCATT TTATTTTAGC TCAAAATATA AAAATATTCA TCAGTTAGCC AAGCTTTTCN GATGAGGAT CATAGCCTCC TCTTTGATAG GGGGTTTCTT GGGTTTCCTT GATTTCATGT TTCAGAGTTT TT

SEQ ID NO:243: (Length of Sequence = 256 Mucleotides)

CTCACACATT CATACCCAAG GAAGAGGCAA ACACACTCAA GTCCAGAGTT CCCAGTGGTG CCGCCCAGAC CTACTGTCCC
GGGGGTGTTA TGGCTGTCCC TCGGCTTCCC CAGAGCAGCC AGGACAGCCT GCACCGNCTN CCAGACTCTC GCAGGAAGGG
GAGCTCTGCC CTGGGGAGGA AACTNACAGG CTGGGAGACA AGACTCCCAT CGCAGGGACA TGCACAGCAG CAGCCACAGC
CCCGGGGAGG GGGCAT

SEO ID NO:244: (Length of Sequence = 220 Nucleotides)

CAAATGGCAG TTCTCGAGAA TCGACGAGGA ACTTAAATCT GGACTCAGGG TTTCAGTGCG GTCTCCGACT CCCACCACCC CGCCCCTCCG NCTGTCTCGC CGCCAGGNGT GACCTCCACG CGAAGGAATC TTCTTCGGAT GGGTGCACCT TGCCAANAGG TGTGGCACCT GGNGGACTAG GAGGCGCCTC CANACTAAGG GCGCTCANTG CGGCGTTCTT

. .

- <u>:</u>
- SEQ ID NO:245: (Length of Sequence = 239 Nucleotides)

TRATECTCA TETAACCTIC TTAATAGIEC CITETCIECT GEGITTETAG CITETAGAGT TETECAAACT GECCETATAA
AAATATTEAT GETETCEATT AAAATGAATE TETETCIECT ACTEAGTETE TETETCIECT TETETCIECT
CETECCATET GIGIGICIET CICTACICCT CIGATTITEN CETETCTETE TATTETEGETA CICTCICCICC TETECCICCG

SEO ID NO:246: (Length of Sequence = 269 Nucleotides)

GGITTCACCA GCGITTAATG TGCTCTGATG TTGACCGTCC CTCTNAGINT TCTGGGGAGG AGGGGGTGGG GCGGAGGGTC
AGGAAAGCAG GCTCAGCTTC CAGGGTCAGG GAGTTGTGGG CCCAGAGGGG CTGTCACAGT GGATGCACCC TGCCCCCTCC
CTCGCCAGAC CCGAGGGTAG GGCAGAGGCA CCTCCTCGNC AGCCTNTGGG CTGCACCCAC AGGGAATNGA GGGGAGGGGC
ACCATTACCA CTGGACCCAC CAAAGACCC

SEO ID NO:247: (Length of Sequence = 297 Nucleotides)

CTATTCAAAG TITACTGACC TCCCCAGCCA GGCAGGCCAA CCCTTCCGAG CAGGGGAAAT GTCCATCTAG CTGCCCTCTG CTGGGTTGCA GCCTATGCCA TGAGAGGGTA CTGGAAGCAG GAGGGAGCCC TGGCTAGGGC AGGCCTTAAA CGCAAGGGAA GCTGAGCAGA GATCTGCACA CTCAACCCCA TITGATATTC TTCTCCTCCT CAGTCATGGC CAGCGTGTTG GTGACTAGAC CGGTGCCCAAT AGTCCGGTTG CCATCTCGCA GGGTGAAAAG ATGGCCTTTC TCTTAAG

SEO ID NO:248: (Length of Sequence = 281 Nucleotides)

ACAACAAGCA CACCAACTAT ACCATGGAGC ACATCCGCGT GGGCTGGGAG CAGCTGCTCA CCACCATTGC CCGCACCATC
AACGAGGTGG AGAACCAGAT CCTCACCCGC GAGGCCAAGG GCATCAGCCA GGAGCAGATG CAGGAGTTCC GGGCGTCCTT
CAACCACTTC GACAAGGATC ATGGCGGGGC GCTGGGGCCC GAGGAGTTCA AGGCCTGCCT CATCAGCCTG GGCTACGACG
TGGAGANCGA CCGGCAGGGT GAGGNCGAAG TTCAACCGCA T

SEO ID NO:249: (Length of Sequence = 383 Nucleotides)

AGGCATCCA CACCGGGGAG CGCCCTACC CCTGCTCCTA CTGTGGCAGG AGCTTCCGCT ACAAACAGAC ACTCAAGGAC CACCTCCGTT CAGGCCACAA TGGAGGCTGT GGGGGTGATA GTGACCCATC AGGTCAGCCA CCCAACCCAC CAGGTCCCCT CATAACTGGG CTTGAAACTT CTGGCCTGGG TGTCAACACT GAAGGTCTAG AGACCAACCA GTGGTATTGG GGAAGGGAGT CGAGGGGAG TTTTGTAAAT CCAAATCTCT GTGGVTTCAT GCTTTGTATA TGCTCACAGC AGGGCACAAT AATCCAAGAG AAGGTCTGTG AGCCCCATT CCAACACCAC AGTAATTATA ATCTTGGCAC ATCAATGGAA TIT

SEO ID NO:250: (Length of Sequence = 397 Nucleotides)

GIATCCIACG TIACAACAAT AATATCATGG GAGAAATAGA AATAGCCTAG TITGCITCCA ATAGAAACTG CITITAACAT
GGGCTGIATA TAAAAATATT AAAGAGAAC AAAACTGTAC ATTTCCTCAT TGCTCCGCTA CAGACAACCC ATGTCATAAC
CTTGTTGCAA ATATTTTTCT CCTATAGCAG TAAGTACAGC ATTAGAAGGT GATTAGAGAG TCTGTTGATG AAACACAAAT
GTATGTTTT ATTGATTTTT ACTTTAGAAC ACTACAGAGT TCCTGGGACC GGGGTGAANG GCATTTAGCT GGGGTGGTTT
GTGTGGGGGT TAAATACCTT CCCACTTGCA AGTGACTTGC CTGTNCCCGC TGCGGGAATC CTGTTNCTTG GGTGGGA

SEO ID NO:251: (Length of Sequence = 276 Nucleotides)

GGCCATAAAA GAAAGAGCCT GTTACCTATC CATAAACCCC CAAAAGGATG AGACGCTAGA GACAGAGAAA GCTCAGTACT
ACCTGCCTGA TGGCAGCACC ATTGAGATTG GTCCTNCCCG ATTCCGGGCNC CCTGAGTTGC TCTTCAGGAC NGATTTGATT
GGAGGGCNGA GTNAAGGCAT CCACGAGGTC CTGGTGTTCG CCATTCAGAA GTCANGACAT GGACCTGCGG CGCACGCTTT
TCTCTAACAT TGTCCTCTCA GGGAGGGNTC TACCCT

.1.

SEO ID NO:252: (Length of Sequence = 314 Nucleotides)

CCTGAACAGT CTGTTCATT TGACTGTTTG GGGGTCTCCC AGTTTAAGCA AGATATTTAA GCCTTATTTC TCTTGGCATG
CTTGGATTCC CCAGTAAAAA AAACTCCTGC CCTGGGCTGA CAATCAAAGT TCTGGGAACT AATATGGATA AGCAAGCTGG
AAATGGAGAA GGCTATTCAC TGTGCCTGGG TCCTACTGTT TTCTGGNTGG GAACTGCTTT TCCATTAGGC CTGGTGTGCC
CTGGAAGGGA MGAGCCTCTT GCAGAGACTA CAATCTTGGA TGGGTCCTTT GCCAAGTTTG AAGGTAGGAA CCCA

SEO ID NO:253: (Length of Sequence = 293 Nucleotides)

GAACACTCTG CTCCAGCCAA GGTGGTGAGG GCAGCTGTTC CTAAACAGCG CAAAGGCAGC AAGCCACAGT CCCACAAGCC
TCAGCCTACC CGTAAACTGC CACCCAAGAA GGACATGAAG GAACAGGAGA AAGGAGAAGG GAGTGATAGT AAGGAGAGTC
CAAAAACCAA ATCAGATGAA TCAGGGGAGG AAAAGAATGG AGATGAGGAT TGCCAGCGAG GCGGCCAGTA GAAGAAAGGA
AACAANCACA AGTGGGTTCC ATTACAAATA GACATGAAGC CTGAAGTGCC CAG

SEQ ID NO:254: (Length of Sequence = 413 Nucleotides)

CITTITCTIA ATATATAAT ATTIACCAAG GCAAGACAGI GATTIATGGA CATITAAATT AGITIAGCIT TGITCTGCTG
TTCTAAAACA TTGTGIACTG TCTGATAGAC TITTAAAAAA CAGIGCTITT CCAGGATGAT TIATGATATG CAGIATTGIT
TATAGATGCC CATGGCTTAA CCTTGAAAAG TCAATTAAGI GACACAATTA AGAGAGATAT GAATAGIGGI AGAAAAAGCA
TGIACTCTGG ATAAGIGGGG GIAAATCTAG TATTTGTTAT TCCTGTCAGI AATATTGTCA NIAGIATTTT TTAGAAGGIT
TAATTTTTTT ATGGGTTATA AATTCATGCC ACTCTTCTGC AATGGGTACC ATCAGTGGGA ATGCNGGAAT TATCCATGCT
TTGGGGGTTA AAA

SEO ID NO:255: (Length of Sequence = 376 Nucleotides)

GGGICCAGGG GAGAATCAAT ATATCIAGIA TAGITTATAT TIGIACCITC TCICCITAAG AGITACAGIG AGIGACICIA
CICCICAAAT GGAGCACCIC TCICCAGGAG AGIAAGAAGA TCACATAAAT AGAAAGIGAG CITIGGACIC TAACAGACAT
AGGITCATAT TCAACICIGC TACITAATAT CCATATIGGT TIGAGITATT TAACCITGAC AATCCACACT GIAAAATGGG
TAAATAATAA ATACCCICCI CICAGAAGIG TIACAAAGIT TATATGAAAT AATGIGCITA AAAAGCIGGG TACATAGIAG
GAGCITAGIC ATTGITTATT TICICCCICA TACCCATACA TGYITCATIC CIACIG

SEO ID NO:256: (Length of Sequence = 241 Nucleotides)

SEQ ID NO:257: (Length of Sequence = 406 Nucleotides)

CAAGGGIGIC CTTCGCCAGA TCACTGITAA TGATTIGCCT GIGGGACGCT CCGIGGATGA GGCTCTGCGG CTGGICCGAT
TAAGAAAACC AAGAGAGGCC GGGCACGGIG ACTCACGCCT GIAATCCCAG CACTITGGGA GGCCGAGGIG GCGGATCATG
AGGICAGGAG ATTGAGACCA TCCIGGCIAA CACAGIGAAA CCCCGGICTCT ACTAAAAATA CAAAAAAATT AGCTGGGCAT
GGIGGCACGC GATTGIAGIC CCAGCTACTA GAGAGGCTAA GGCAGGIGAA TCGCTTGAAT CCAGGAGGIG GGGGTTTCAA
TGAGACCCGAG ATCGIACCAC TGCACTCCAG CCTGGGGCAA CAGAGTANGA CTTCGIAACC CCCAACCAAC CCNCCAACCC
CCCGCC

SEO ID NO:258: (Length of Sequence = 157 Nucleotides)

GAAAAGAAGG AAGGAAAGAG GCGACCGACG GAGGAAACGA GAGAAGGACG GAAAAGAAGGA GAAAATGCTG GAGCAAAGGA GGTTGGTTAC ATGATTTCTC TAATGGCAAT GAGCTGCTTT CTGGATGAAA TACAGAATCA GAGCGAGACT CCGTCTC

SEO ID NO:259: (Length of Sequence = 361 Nucleotides)

AAGCAGATAT AAATGGGACC ACTGIGAATC AAAGGGGAAA AATTCCAGGA AAAAAAAATT CCAATAGCTT CACAGTTTAA
CTGAGGTTTT GGAAAAACTT AAGTGAATTC AGCTGATGTT TGAAATATCT GTCTACATTT AATTAGATGT GTTGTATTTA
CCAAGGAGGC ACAAATATGT AGTTCTGTAG ATTTTAATAC TAACTTTTCC AGTAAGAAAA ATAATACCAG GTGATTTCAA
AAAGGGCAGT GATCTATAAA CACTCAAAAT GCATCTTTGA ACAGGGGAGC AGAAATAGCT AATTTAATGA AAACAAACCT
TAAGCACTTT ACTTGGCTTC TAATAAGGCA TCCCAAGAAA A

SEO ID NO:260: (Length of Sequence = 349 Nucleotides)

CAATACATGT ATACAGTGTA CACTGATCAA ATAAGAGTAA TTAGCATATT TATCACCTCA TITCTTTTGT GGTGAGAACA
TTTAAAATCC TITCTTTTTG CTATTTTGAA ATATACAGTA CATTGCTATT AAGTATAGTC ATCTGGCTGT GCAATAAAAC
ACCAGNACTT ACCCCTCCTG TCTGTGACTT TGTACCCTGT TCACCACCCC TCCAATCCTC TAGTAACTAC CATTCTACTC
TCTACTTCTA TGAGCCTGAC TTTTTAAAAT TCCACATGTA AGTGAGATTA CATGGTATTA TTCTCTCNGT GGCTGGCTTA
TTTCACTTTA ACATAATGTC CTCTAAATT

SEO ID NO:261: (Length of Sequence = 415 Nucleotides)

GGAAGATGAG GATCTAGGTG TGAGCGTGCA GAGCCCTGAG GCTGGGCAGG CAGGGAGCTC TGCCTGCACA ATGATGTAGC CATGTGTGGC CACACCAGCA CTGGGCAGCA CCTCTGGGGA GGGGGCCAGG GCAAGGACAA CTGGAGAGAC AAAGCCAGAT GGGGCCACGT CCTTAGAAGT GTGTGTGTGT GTGTGTGTGT GTGTAATACG CAGGCCAGAA ACACACCATG TAGGTCAGGC AGGACAGAAA CACACCATG TAGGTCAGGC AGGACAGAAA CACACCATG AGGTCAGGC AGGACAGAAA CACACCATG AGGTCAGGC ATCACCTGAG GTCAGGAGTT CGAGACCAGC CTGGCCAACA TTGCAAAACC TCATCTCTAC TAAAATTCTA AAATTAGCCA GGCGT

SEO ID NO:262: (Length of Sequence = 382 Nucleotides)

GECATGGGT CIGGCTTIAA TGIGIAACIG ACGIGGGTCA CIGAAACIGT TCAGGCIGAT CITGAACICC TAGGCTCAAG
TGATCCIGCT GCCTIGGCCT CCCAAAGIGC TGGAATIACA GGAATGAGIC ACAGCACCCA GCCGGCTGIG TTITGITTIT
TGITTITIAC CCCGACAGGI NCICAGICAG TCGTIAGCIG GAGIGAAGIG GCGIAACACA GCTCACTGCA GCCTIGATCT
CCTGGGCTCA AGIGATCCIT CCATTTCITC CITCCAGAGI AACTGGIACT GCAGGCCCAC GGCACCACAC ATGGCTAATT
TTTAAATTTC GIAGAGACGA GGICTIGCCA TGITTGCTCA GGCTCCAGCT GTTGTATTCT TT

SEO ID NO:263: (Length of Sequence = 447 Nucleotides)

TGTATCAACT CAGAATTTCC AGAGAGCTCT TCCTGGCTGA AAAGATGTCC AAGGATCATC TCCGGAATGG AAGAGGTGAG
GCCTGTTAGC TTGTGGGCTG CCCAATCCAT CCAACCCTTG GCATTGGGAT CAATGTTGAT GAGGACAAGA CCTTCAACAG
TGTCCGGGTG GTTAAGAGCA TATCTCGCCA GGATGTAGGC TCCAGCTCCA ACACCAACTC CAATTATTGT AGAGAAAATTT
AGGTACTGCA GGACGCAAGG GATCATGTCT GCAAGCTGGT CCAGAGATGG GTACTGATAT CCCAAAGGGA ACACAGGGC
TCCCTCTTCC ATTCCAGGGG CATCCACATG GACCCGCACA AAGTTCTGAA TGATTTCCTG CATGTCCTCG AACTKGAACA
GTGGCTGGAG GAAAGATTTA TAGTTGAGTC CACATCGGGT AGGTAAG

SEC ID NO:264: (Length of Sequence = 317 Nucleotides)

TTTTCGCTGT CAACAGACAG TTTATTCTAT ATACAAACAC AATTTTGTAC ACTGCAATTA AATAGAATGG AATGAGCGCT
CCTCCGCATT CCTCCCCGAG TGACTGGTTT GGCCGCCGGC CACTCCATCC CCGAGTGGGA CTGGACCACG GCCCTGGNTG
CTGCCACTGA TGTTGGNGCC TGCACCCCAC GTCCCTATGC CCGAGGCGCA ANTCTGCTCT CCCGGGGACC CCAAGNCTGG
NGCACACGCG GGGAGGGCGG GGCCATGGAG AAGGCACTGC AGGGAGCACC AGGCAGAGCC GTGTTGAGGC CGGCCCG

<u>SEO ID NO:265:</u> (Length of Sequence = 270 Nucleotides)

GCAGAGCAGG TGGAAGTGAT CAGGAACCAT AGTTGACAGT TCCAATCAGT AGCTTAAGAA AAAACCGTGT TTGTCTCTTC
TGGAATGGTT AGAAGTGAGG GAGTTTGCCC CGTTCTGTTT GTAGAGTCTC ATAGTTGGAC TTTCTAGCAT ATATTGTGTCC
ATTTCCTTAT GCTGTAAAAG CAAGTCCTGC AACCAAACTC CCATCAGCCC AATCCCTGAT CCCTGATCCC TTCCACCTGC
TCTGCTGATG ACCCCCCCAG CTTCACTTCT

SEO ID NO:266: (Length of Sequence = 297 Nucleotides)

ATEAGGCGAG GCCTGCGAAG TGGCTGGCAT GCAGCAGGTG CTAATGAGTG TTGCAAAGGT GATGTCACGC AGGCAGCTTC
CCGTGGCCAG AGAAACATTG CAGAGAAGGG ATAAGTAGGG CTTAGTGACT TTGACGGGTC AATGGAAGAA TGACCCAAAG
AAGGCTTCAA GGCCAGGCCT GCAGTTCTCC ACCACAAAGG CCCTCACTGA TAGCACCCAC TCCCCCACAC TCAGCTTTNG
GGCCTAGGTC TGGGTCACCC AGCTAGAAGC CACAGGACCC TGAGGCGTCC GAGGGGT

SEQ ID NO:267: (Length of Sequence = 387 Nucleotides)

CTTGITITCA TCATGAGCIC GATCAGATGI CICICGATCI TCAGACIGGI GGIGICCIAI AATGICCIGI GCACGCATTC
TTGAGCITIC CAGGATTICT GICTGITCIC TCIGITTATC TACAGAAGAA ACTITCICCI TGAGTICCIG TICITCGIAG
CGCCTTGAAC TCICITTCCT TICTGGITTA CGATCCICCT CTTTCCATCT ACCCTGTCTG TCTTCTGTGA GGIGCGAGGG
ACTAAGAGAA CGAGATTCTT GAGGICGIAC AACTTGGCTC AAGAGTCTGT GTTTTTTCAT TININATCAT CTCCACTGTT
GTAGGCATCA CTGICCGGAG AATGITCACG CCGGGGCTTT CGGGGGACTG TCTAGGGCTG GGACTCC

SEO ID NO:268: (Length of Sequence = 318 Nucleotides)

CCTGAAGGIT ACCICTITGG AGAGAACATG GATCIGAACT TCCTGGGCAG CCGCCCGGTC CAGTITCCCT ACGTCACTCC
TGCCCCCCAC GAGCCCGTGA AGACGCTGCG GAGCTGGTGA ACATCCGCAA AGACTCCCTG CGGCTGGTGA GGTACAAAGA
CGATGCCGAC AGCCCCACCG AGGACGCCGA CAAGCCCCGG GTGCTCTACA GCCTGGAGTT CACCTTCGAC GCCGATGCCC
GCGTGGCCAT CACCATCTAC TTCCAGGCAT CGGAGGAGTT CCTGAACGC AGGGCAGTAT ACAGCCCCAA GAGCCCCT

SEO ID NO:269: (Length of Sequence = 422 Nucleotides)

ACAGGICIAT TCAGGICITT TGCCCATTIT GAAATAGCAT TGCTTGTTCT TTTGCTGGAT ATTAACCCCT TGTCAGGIGC ACAGGITTGCA AGTTACCTT TCTCATCCTA TAGGITTATCT CCTCACTCTT GATTGTTTCT GITGCTGTGC AGTAGCTTTT AAGTTTGGGT TAATACCATT GIGTTTTCTC TGCTGCCCCTT TTAAGTTTCA CTGGGTCAAA AGTTTAAAAT TTGTGAATTC CTATATTTTT AGGGCAATTC TCCTGCCACT GITGGAATTA TGCCTCAATC TATGCAGTAG AATATTAGTG TGAAATGCTT CTGTACCAAT GGAGATGATG CTGGATGGTC TCTATCATAA ACCCATACCT CATCAACACA AACTGCAATT ACACAAGGGC TCTATATCAT GGATCTCCAT TT

SEO ID NO:270: (Length of Sequence = 376 Nucleotides)

GAAGAAGAC CCAGACCTAG GGGAGTATGA TCCACTTACC CAGGCTGACN, CHGALGAGAG GGAAGACGAT CTGGTGCTTA.

ACCTGCAGAA GAATGGAGGG GTCAAAAATG GGAAGAGTCC TTTGGGAGAA GCGCCAGAAC CCGACTCAGA TACTGAGGTT
GCAGAGGCTG CAAAGCACAT CTTTCAGAAG TCACCACGGA GGGCTACCCC TCAGAACCCC TTNGGGGCCT GGAACAGAAG

GOGGCCTCCT CCCTGGTGTC ATATGTGCGC ACGICTGTCT TCCTGCTTGA CTTTGGGGAT CTCGATGATC CTGGTGCTCC
TGTGTGCTTT CCTGATCCCC TGTCCTCCCA GAGATCTTGA CAGAACTGGA GCCGCA

SEQ ID NO:271: (Length of Sequence = 346 Nucleotides)

TGITCACGIT CCCTTTCTIT GICTITCTIT TICCIATCIT TATCIATACI TCGACICCTC TCCTTTTTCC TCTCTTGITC
TTTAGCCTCA CCTTTATGCT TATGACTGIN CCCACTAAGA TITCCACGIT GATCATCAAT TITACGACTCA TCTCGACTCC
TACTGCGACT GCCACGATTG GITCGTCTAT CCCTTGAGGG ACTTCTACGA ATGCTTATGA AAAAGAATCA AGTTGCACCA
CAAATGITTC ATAGCAGTAG GAAATTTCTT TTAGAGACTT CTGATGGGAA ATTTGAAGTG TATGTTGCTA TCAGATCAAG
TGCAGGAGAG GTATAAGGCT ACTGGA

SEO ID NO:272: (Length of Sequence = 394 Nucleotides)

GTIGITIGITG TIGAGICGGA GICTCGCACT GITGCCTGGG CTGGAGIGCA ATGGIGCAAT CTCGGCTCAC TGTAACCTCC
GCCICCCAGG TICAAGCCAT TCTCTTGCTT CAGCCTCCTA GTAGCTGGGA TTACAGGCAC CTGCCAGCAC ACCTGGCTAA
TTTTTTTATAT TTTNAGTACA GACAGGGTTT CACTATGTTG GCCAGGCTGG NCTTGAACTC CTGACCTTGT GATCTGCCCA
CCTCAGCCTN CCAAAGTTTT TCAGAATTTT TTAAGGAAAC ACTTTTAACC CTTAAGGCTT TCTTTCAAAC TCAGATCCCC
TTACACAATT GATCAGACGT GGCAAAGTTT TGCTTCAAAG TTTTTGGACT GGGTTTCCAC TTTAGGCTTA CTGA

SEO ID NO:273: (Length of Sequence = 259 Nucleotides)

CAACCIGIAC CCAGGCIGCG AGAACGIRAG TITRAGGAGC CGCAGCAIGA TGITCGAGCC GGGICITACC AAAGGRATGC TGGAGGIGIT TKIGGCCCCG ACCCACCACC CGCACIGCIC GGCCGAIGAC CAGICCACCA AGGSCAICGA CAICCAGAAC GCITATITRA AIGGAGIIGG CGAITICAGC GIGIGGGAGI TCICIGGAAA TCCIGIGIAT TICIGCIGIW AIRACTATIT TGCIGCAAAT AAICCCACG

SEO ID NO:274: (Length of Sequence =348 Nucleotides)

TCCCAGTTGT CCCGATTGTA ACTCAAAGGG TGGAATATCA AGGTCGTTTT TTTCATTCCA TGTGCCCAGT TAATCTTGCT
TTCTTTGTTT GGCTGGGATA GAGGGGTCAA GTTATTAATT TCTTCACAGC TACCCTCCTT TTTTTCCCTA TCACTGAAGC
TTTTTTAGTGC ATTAGTGGGG AGGAGGGTGG GGAGACATAA CCACTGCTTC CATTTAATGG GGTGCACCTG TCCAATAGGC
GTAGTATCCG GACAGAGCAC GTTTGCAGAA GGGGGACTCT TCTTCCAGGT AGCTGAAAGG GGGAAGACCT GACGTACTCT
GGGTTAGGTT AGGACTTGCC CTCGTGGT

SEO ID NO:275: (Length of Sequence = 396 Nucleotides)

GITTIGGIGAA TITIGGICTIGI GATAAAATTIG GAGITCAAGA AACAAACAGG AAACTACAAG TGCCCCTTCG CCCCCAGGTC ACCGGAGTIGG CAGGGCAGTIG ACCGCTGCTC TCAGGCTGCC CAGTGTGGAC CTGCCTGTCG GAATGCTCCT CCTCCACGTC CCCTCGCTCC TGTGTCCCCAG CCACATGCAC CTTCCCTCTA CCTCTGGGAT CCCTGCACCA GGTCTGCCCC TGTCTTCTCA GGGCTGCTCC TNTTGGNCCA CAGGACCTCA GCTGGAATGT TGCCTCCTCC AAGAGGCCTT CCTGACTATT CAGCTCACAG TGGCCACCCA GCCACAATCT GCCATGTGCT TTGGGGGATT GTCTGTTAAC TGGCAACATA CTGGCAGCCC ATAACT

SEO ID NO:276: (Length of Sequence = 381 Nucleotides)

GETGTCGGGG AGGCTGCGCA AGGGGGGGAG CCCGGGCAGC CGGCGCAACC CCCGNCCCAG CCGCACCCAC CGCCGCCCCA GCAGCAGCAC AAGGAAGAGA TGGCGGCCGA GGCTCGGGAAA GCCGGTGCGT CCCCCATGGA CGACGGGTTT NTGAGCCTGC ACTCGCCCTC CTATGTCCTG TACAGGGACA GAGCAGAATG GGCTGATATA GATCCGGTGC CGCAGAATGA TGGCCCCAAT CCCGTGGTCC AGATCATTTA TAGTGACAAA TTTTAGAGAT GTTTATGATT ACTTCCGAGC TGGTCCTGCA GCGTTGATGA AAGAAGTGAA CGAGCTTTTA AGTTAACCCG GGATTGCTAT TNAGTTAAAT GCAAGCCAAT T

SEO ID NO:277: (Length of Sequence = 206 Nucleotides)

TTAATACGAC AGGCCTGGCG CCCGAGTAAT TCAAGCCCTT CGGAAGTGTC ACCGGCTGCC AGGCCTCGGA TGCAATCCTG GAGGCGGGAG ATTCGGCCTN AAGACTGGCT CGAGCCGCCC AGGGGCTCCA TGGGAGACTA ACGCGGAAGT YCCAGCCGTC CCAGTGCCGT GACGTCCCCC CTTGGTGGGG CCTGCACCCG ACTACT

SEO ID NO:278: (Length of Sequence = 260 Nucleotides)

ACCTGTAATC CONGCACTIT GGGAGGCTGA GGTGGGCAGA TCACGAGGTC AGGAGATAGA GACCATCCTG GCTAACACGG TGAAACCCCA TCTCTACTAG AAAAATACAA AAAATTAGCC GGGCATGGTG GCGGGCGCCT GTAGTCCCAG CTACTCGGGA GGCTGAGGCA GGAGAATGGC GGGAACCCGG GAGGCGGANT TGCAGTGAGC TGAGATGCGC CCGTCTCTCC AGCCTGGGCA ATAGAGTGGG ACTCCATCTC

SEO ID NO:279: (Length of Sequence = 308 Nucleotides)

GIGICIGGEC TCAGGGITGG CCAGCITGCA GAGGAGCAAG CIAGIAGAAA TATIGCAGGG TICCCAAAAC CAGGICAAGC
AAGATGCCAT GICACCCCTG AGCATGCCIG TCITCCCAGG GGIGIACCTC TIGGCTGGCA AAGCCAAGGC CAGIGGGNAC
TTGIATAAAT CACATGGGIA TGITCITGGI TCAGIGATCI TGGAGIGATG ATGGIAACIN ATGAACAGAG AACTITYYAG
AACTIKGGIC CIGICTTCCI CCCIGAACCT AGACAAGTIT CACCCCTCCI CCTGIACCCA ACCCCATT

SEO ID NO:280: (Length of Sequence = 402 Nucleotides)

ATTITAGCAG CITICITGAA ATTITAAAATA TATGIGIAAG TATCICATIT ATATGCATIT CIAGTITCIT TATACAACAG
AATAACTICI TITACATCAA ATTICIGAAT TIGACIAAAT TIAGAAATAA TGGAATCICA TCCATIAAAT ATAGICATAG
AAGGAAGGAA ATATGAAAAT TAGGATTICA GATGITTGAA CATAAAAGAT AATTITAAAC ATTGICAGTA ATCIATTICT
TITTITITIC GAGACGGAGT TITGCICTGT CACCCAGGCT GGAGIGCAGT GGCGCGGTCT TGGCTTACTG CACCCTCTGC
CICCCAGTIC AAGIGGATIC TCCTGCCTCG NCCTCCTGAG TAGCTGGGGT TACAGGGGCA TGCCAACATG CCGGGGCTAA
TT

SEQ ID NO:281: (Length of Sequence = 313 Nucleotides)

GAGAATCOGT CITAAAAAGA AAAAAAGAAA ATTATAGAGG GAGATGAGGT GGGACAGAGT CTGGCAGTTC ATCAGGGGGA CTGAGAAGGT GGCATTTGGA GGAGAGGAGG CAGTGAGCTG TGCAGTGTCC AGGCAGCCAC CCTTCCCCAGC GGCCACCATG ACCGTGTCCT CATTGCTTTA ACCATTAGTA ATCATTCATT CATTCATTCA TITATCCGAC GTCAGCTGGA GGNCCTGCCC GNGGGGCATG CGCTTAGATT TNGGAGGCCT TCCGGGATGC TTGCGCTCCA ACCGGGGAAG GCCGACTTGG GCT

SEO ID NO:282: (Length of Sequence = 217 Nucleotides)

TEACCTCAGT TGATCCACCC ACCTTGGCCT CCCAAAGTGC TAGTATTATG GGOGTGAACC ACCATGNCCA GCCGAAAAGC
TTTTGAGGGG CTGACTTCAA ATCCATGTAG GGAAGTAAAA TGGANGGAAA TTGGGGTGCA TTTTCTAAGG ACCTTTCTAA
CANATGGCTA TAATNTAAGG GGTTTAGGGT CCTTTTTTTT TTTTCAGGGA TACATTT

SEO ID NO:283: (Length of Sequence = 327 Nucleotides)

TAGAGAGGCC TITACTCCTG GICCCATGGC GIAAAGATGI GGCTGGGCCI GACAAGGCTC AGCCTCCAGI CTTAAGATGG GCACAGAAGG GCAAGAAGTA AGATGACGAG TCCCAGAATT AGGACAAGCC ATGAGCCAAG GCCTGGTCTG AGCAAGGGCA GCCCCCTGTC CCAGACACAG GCACCCCCAA TCTCACTTTG GACAGAGCCA ACGTGGGGG ATCCTCCCGG GCCTGGGCCT GTCAAGTCTG CCTGCAGGAC CCTGCCATTG TGCTCAAATC ACAACCATTT TTTGCTTCCA ACATTTTAGG GTGCTTGTGC AGTGAGT

SEO ID NO:284: (Length of Sequence = 340 Nucleotides)

CTITIGGAAAT GIAAATTGIT ACAAACITAC TITIAGAGCAA ATTIAGICAT CCITICAAAAA TITIAAATGIA TACITATITIC CTAAGAATTC GITIGGCTCA CACAATTGIG AAAAGATAGA TGIACACCAG TGITCATIAC AACAATTATG CAACAAATCT ATTIATGIGCC AGACATTATT CGGAACICTG GGAATACATA AGIGAACAAA GCAGATTCCT GATCICAGGA CCTGGGGICA GGGGICAGGA GAAGCCAAAA AACACGCING AGAAATACTI TATGCAGTGT GGGGGGAGTG CTACCAGCAG AGCAGGGGAT GGRGATGIGA AATCITGIGT

SEO ID NO:285: (Length of Sequence = 335 Nucleotides)

GACATICACG GAGGIGGGIT CGACCICCGG TICCCCCACC ATGACAATGA GCIGGCACAG TCGGAGGCCT ACTITGAAAA
CGACTGCTGG GTCAGGIACT TCCTGCACAC AGGCCACCTG ACCATTGCAG GCTGCAAAAT GTCAAAGTCA CTAAAAAAACT
TCATCACCAT TAAAGATGCC TTGAAAAAAGC ACTCAGCACG GCAGTTGCGG CTGGCCTTCC TCATGCACTC GTGGAAGGAC
ACCCTGGACT ACTCCAGCAA CACCATGGAG TCAGCGCTTC AATATGAGAA GTTCTTGAAT GAGTTTTTCT TTAAATGTGA
AAGATATCCT TCGCG

SEO ID NO:286: (Length of Sequence = 399 Nucleotides)

GCACAATTAT TAAAAAGAGG CCACTTAAAT TCAACTCTCC ATGGATACAG TGTCTGTGGC AATGTTTAAT TAGAGATTAA
AATTGAGGAA TTGAATAATT GAGGTTGCTA ATGAATTTGA AAACTCAGCA AAGCAAGGAG AGCTGAGCGT TTTTCCGACT
TAGCTTTTCT TTCTCTAACC CTTTTCTCAT TTCCTACTAT TATCACATNT CTGGCCTTGA CTGCTGAGGTT TATTACTACC
CATAACCCTG GCCTAAGTGG AAACAAAAAA GCTGTAGCCT CTTTGCTGAG CTCCTGGAGA CATTTGGTCT ATTGGATTTA
TGACATGTTC AGAAGCTTGC AGTTGCAGGA GGCTGACAAT GATGAAAATG AGATATGNTG GGCCACCACG CTTTTCTGT

SEO ID NO:287: (Length of Sequence = 294 Nucleotides)

TTCCAGTIGA ATTCACCAGI GGACAAAATG AGGAAAACAG GTGAACAAGC TTTTTCTGIA TTTACATACA AAGTCAGATC AGTTATGGGA CAATAGTATI GAATAGATIT CAGCITITATG CTGGAGTAAC TGGCATGIGA GCAAACTGIG TTGGCGTGGG GGIGGAGGGG TGAGGTGTG GCAGACCTTT TTTTAAGATT TTNCAGGTAC CCCTCACTAA AGGCACCGAA GCTTAAAGTA GGACAACCAT GGAGCCTTCC TGTGGCAGGA GAGACAACAA AGCGCTATTA TCCT

SEQ ID NO:288: (Length of Sequence = 391 Nucleotides)

TCTACAGATG AGGAAAGCAA GCCTCAAGCA AGGGGGGCCT GATCCTTTCC CTGTTCCCTG TGTATTCCCT GTCTGTGGCA AAGCCCATTG CCTTGATTCT CTTCTCTTTA CTTTCATGTT GAGAAAGTAGT TTCTTTCTGC AGTTTATTTA ATTTACTGGC AAAATGACGT ATTTTTTTTT CAGCAATGTT TCAGCTAGAT ATTTGCTTTA TGCATGTAAT GTCAATGAAG TACTCATAAG TTTTCAAGAA ATGACTGATA TAAATCATGT GTTCACTAC ATAGTCTAAA TATTTAGTAT TTGGTCATCT ATTTTAATAT GTTCAAAATTC TGTTAAACAA GNCATAGTCA CTATGTGAAG ATAAAAATAG NCAAAGTTGC ATTATGACTT T

SEO ID NO:289: (Length of Sequence = 198 Nucleotides)

CTTATATTCT ACTITATTIC GIAAAACTCA GAAACTAACA ATTCACATCC TCCCACCTTC TTCTTTCCGA AGAACGCAGT
TTGCAGAGAC AAAAGGGCTG TGGCGTGGGG ATCATCCACC ATCTCCAGGT TTTACACCCA GGCTACCCAT GGCTTGGCAG
ICAUGCCTCT AUGCTGATTG CTCTCAGAGG CAATAGAA

SEO ID NO:290: (Length of Sequence = 353 Nucleotides)

GETTITCATO TICGGITTAC AAAAGICCIA CIATITATIT ATTITAACIT TAATITAAAT ATCACCIACO TIAGGIAGAA
GITTICCITT GIGIAATATA ATATAAAACO GACATITCIT GGGGCATAA TAGTAAAGAT GITAACATIT TITGGITCIT
TITGGATGCT GIATITGIGC TICTICIGAA AGIGATGIGI GCCAAGATGG CICATGIAAC CCAGTITTGA CIAGGCTATT
GATATICIGI CIGGITAATI TATIGAACIG GCITAAAGCT ATACATATIT CCTITITAGNIGIAA GATATICIAG
ATATATIGGT CIACIGATIC ATAATATCAC TGG

SEQ ID NO:291: (Length of Sequence = 163 Nucleotides)

CCTGGTAGGC CTGCTACACA GTCTTGCAAC GNCCCTCGTG CTTGGGCTTC TGCGGTGAGG CAGGGGAGTC TGCTTGTCTT
AGATGTTGGT GGTGCAGTCC CAGGACCAAG CTTAAGGAGA GGAGAGCATC TGCTCTGAGA CGGATGGAAG GAGAGAGGTT
GAG

SEQ ID NO:292: (Length of Sequence =397 Nucleotides)

ACGGGAAGGT GAGTATGINA GIATGINIGC CAGACAATGG TGTTICCATG TCAATGGAGG TTTCTCAGAG AGAGGTGATC
TGGCTGGAGA AAGCTTAATC TGGTGGCAAT GGACAGGTGA CTTTAAGAAG TGGGGAACGA GGGAAGGAGG CCAGTTTGAA
AATNATAACA AGGGTCCAGA CTCAGTGATG CAGCAGTGAC CATGAGAACA GAGCAGCTGC AGGTAGAAGA TGGAGACAGA
ACTNGGGAGA TCTGGTGGAG GTAAGCCGCG TGGAAAGATG ATGTCAGGTT TATACCTAGA GGACACATGA TCCATTCACA
AAGCCAGGGG NAACCTAAAG AGAAAACACT TAGAATTTIN GGAGAANAGG CTAGGGCTGG GCCTTAGACA TGGGCTG

.SEO ID NO:293: (Length of Sequence = 360 Nucleotides)

GAGGIAAAAT TIACATACAG TGAAATCCAA ATCITAAGIG TACCACTAGA TAAATTTTGA TAAATGCATT ATGCCIGGIC
TTCACACACC CITTTCAATA TATAGAAAAT NICCAGATAA TITATTTTGT TGITTTTTTC ACACACTAAG TICTAGACTT
TTCCAGGICC GAGGGAACTA TTAGGGGGGA AAGTACTTGT NATAGIAAAA AAGATTTTAG GIGIGITTGT TTTTAAGGIG
CAGAAACACA TCGCAGATTT AAGGICTGCA ATCITCTGCTT TTTGTTATTG TTCCAGTTTT GATCTCAGIG ACATTACAAG
CAAGCAGAAA CACTCAGACA TGAAATGGCC CAG

SEO ID NO:294: (Length of Sequence = 321 Nucleotides)

TITITITICAG GNITCAACCE TITIATIGGE AGGITTIGIT TICTGIGAAA TACACIAGAG GGIGGGGAAG GGGACACATT
CACITIGCAA GATAAGGGIT TCCCACCACT AAAGGAAAGG CATGGGGCAG GGCACACTGG GGITTGGGIC CGITTTCCCA
CCTCCTTCTG CTTGGCTCAC TITTCTTTTC TCTCAGCAAG TACCACAGAA CACAAAGACA AGAAACAAAA CAGCAAATCA
ACCTCCAACG GGGCCATGCC AAGCCTTCCC CACTCCCCCA GGCTGGGCAA GGGCTGGGAG GGGCTGGGG CAGCTCACTC
G

SEO ID NO:295: (Length of Sequence = 165 Nucleotides)

GACACAGG GOCTICOGGCC COGCACAGGG GGCATGTCCA GAGGTGCTGT GTGTCACCAA CTGGTCTTCT AATTTGGAAG GAGTTGGAAA GGCCTTTTTG TTGATGAAAA GTTGGAAACA GTGGCACATA TCTNAGAGGG AGGAACGAGG CAGOGTGGTG AAGCG

SEO ID NO:296: (Length of Sequence = 315 Nucleotides)

CHARTACAGG TAGTGCCCAG CTGGTTGCGC TGGCCCAGGA AAATNCTGCT GTGTCAAATA CTGCTGGCCA GGATGAAGCC ACAGCTAAGG CTGTGTTGGA GCCCATTCAG AGCACCAGTC TAATTGGGAC TTTAACCAGG ACATCTGACA GTGAGGTTCC

AGATGTGGAA TCTCGTGAAG ACTTAATTAA AAATCACTAC ATGGCAAGNA TAGTGGAACT TACGTCTCAG TTGCAGCTGG CTGACAGTAA GTCAGTGCAT TTTTATGCCG AGTGCCGAGC ACTGTCTAAA AGACTNGCCT TGGCTGNAAA GTCTA

SEQ ID NO:297: (Length of Sequence = 244 Nucleotides)

AGIACGGI'IN NCGCINAAGC TIGAINATCG RATIGCCAAT CINCATATIT GIGITAGAAT CATITGITIT TGIGICITCA
TGITICIATA AGATAGGACC AATATICITI ATIGGGCITT GATITIATIT TGIAACITAA ATGIATITAAG GCAATAAATG
TAATITITCCA CINAAAACTA TCATTATAGA TTIGGITACT ACCTACIGCT CAGCAATITT TTITCTTATC AAAATICITC
CTGG

SEO ID NO:298: (Length of Sequence = 152 Nucleotides)

CCTGAACAGG TAATGAGAAA AATTTACACA CAAGTGATTT TGAAAACAGA ATGGGTTGCT TACAAATTAC AGGAAATGTT ATAACACAAA CCAGAAGAAT TCAATGGAAG GCAATAAGGGAAAT GAAAATTATA AAAGTATCAN GA

SEO ID NO:299: (Length of Sequence = 374 Nucleotides)

CGATGATTT AATGICATCA CACGITGICT CAAAATGAGI GGIGGCATCA TATGIGCGG AAATAAAGAT CIGGCITTCT
GITCCCAAGI CITTIGGIAC CAGGAGGICA CIGATGCIAA CAAATTTCIG TICAATTGGI TCCAAGAGCI CCAAAGCIGG
TCIGATTICC TICTCAGGCI CCITGGITTC CACAGITGIA CIAACTATAG CAATGIACTI CCCTTGIGCI GCTACATTGI
GCGCAAAGGA GATCATGCAG ACGIAGATAT CIGACTITCG ATTGACTTTG GTICTGIGGA ATAATGATCT GGCAGGAGIT
GGCATCATTG GTGITCTTTG ATGGGGGTGG CIGAGGGATG CAAATAACCT CTTG

SEO ID NO:300: (Length of Sequence = 365 Nucleotides)

GGCTCACCAA GCTCAGCAAG TACGTGTACT TCTTCGAGGC CTGCCGGCTG CTGCAGAAGA TGATTGACAT CTCCCTGGAT
GGCTTCCTGC TGACTCCGGT GCAGAAGATC TGCAAGTACC CTCTGCAGCT GGCCGAGCTG CTCAAATACA CGCACCCCCA
GCACAGGGAC TTCAAGGATG TTGAAGCCGC CTTGCATGCC ATGAAGAACG TGGCCCAGCT CATCAACGAG CGGAAGGGTA
GACTTGAGAA CATCGACAAG ATTGCTCAGT GGCAGAGCTC CATAGAGGAC TGGGAGGGAG AAGGATCTCT TGGTCAGGAG
CTCAGAACTC ATCTACTCGG GGGGAGCTGA CCTCGGGTTA CACAG

SEO ID NO:301: (Length of Sequence = 224 Nucleotides)

GGTATTCAAA CAAATAGCCT GAGAATTING GGGGGATCTG AAATAGAGTA CTATGCTATG TTGGCTAAAA CTGGTGTCCA TCACTACAGT GGCAATANTA TTGAACTGGG CACAGCATGC GGAAAATACT ACAGAGTGTG CACACTGGCT ATCATTGATC CAGGTGACTC TGACATCATT AGAAGCATGC CAGANCAGAC TGGTGAAAAG TAAACCTTTT CACG

SEQ ID NO:302: (Length of Sequence = 363 Nucleotides)

AGTITICACTO TIGITGOCCA GGOTGGAGIG CAATGGOGIG ATCTOGGOTC ASTGCAATCK GCACCITCOG GKTTCAAGCG
ATTCTCCTGC CTCAGCCTCC CAAGTAGITG GGATTACAGG CATGCGCCAC CATGCCCGGC CAATTTIKTA TITITTCGTAC
ACACAGGGIT TCTCCATGIT GGICAGGCTG GICTCAAACT CCCAACCTCG GIGATCCGTC CACCICGGCC TCTCAAAGTG
CTGGGATTAT AGGCATGAGC CACTGTGTCC GGCCAGCTCA AACAATTTTA ATGCTTCTIT CAAGACTATT AGAAACCTTT
AATTGCTTCT TAAGTTTCTC CCCCAACTAT GGAGGAAGCA TAT

SEO ID NO:303: (Length of Sequence = 253 Nucleotides)

ATGCAGGAAS ATCTACCARG CAAATCGAAA ACAAAAAAAG GCAGGGGTTG CAATCCATCT CTCTGATAAA ACAGACTTTA
AACCAACAAR RRTCAAAAGA CACAGAGARG GCCATARCAT AATAGTAAAG CGGATCAATT CAACAAGAAG AGCTAACTAT

CCTANATATA TATGCACCCA ATACAGGAGC AACTAGATTC ATAAAGCAAG TCCTGGAGGT GCCTACAGAG GAGGCTTAGG CTCCCACACA TTA

SEO ID NO:304: (Length of Sequence = 416 Nucleotides)

TTTTTTTEAG ATGGAGTACT OGCICTCTTG CCCGGGCTGG AGTGCAGTGG CGCGATCTCG GCTCACCTGC AACCCCTGCC
TCCCCAGTTC AAGAGGTTCT CCTGCCTCAG CCTCCCGGGT GGCTGGAATT GCAGGCACAC ACCACCATGC CCAGCTGCTT
TCTTGTATTT TTAGTGGAGA CGTGGTTTCA CCATGTTGGC CAGGCTGGTC TTGAGCTCCT GACCTTAAGT GATCCGCCAG
CCTTGGCCTC CCAAAGTGCT GGGATTACAG GCGTGAGCAC CGTGCCCCAGG CTGTTTTTTA ACTGACTTTG GATTTTACTC
CCTTTCTATG CAAATTTATT TTAGAATCTG TTCCTTAACC TTAGGGGGTT GGGTTAGACA AGTTTCAAGG GAGCCTCAAG
TGKAAATTGC TTAAGG

SEQ ID NO:305: (Length of Sequence = 223 Nucleotides)

CACACCCAGC TAATTTTTGT ATTITTAGTA GAGACGGGT TTCACCATGT TGGCTTGGCT GGTCACGAAC TCCTGGCCTT
GAGTGATCCC CCTGCCTCAG CCTCCCAAAG TGCTGGGATT ACAGGTGTGA GTCAGCGTGC CCAGCCCAGA TTTTATTGTT
TTAATTACAA ATTITACGTT AACTGATTCT GCACATTTAT ATTIGCACAC TTGTGCTAGT GAG

SEQ ID NO:306: (Length of Sequence = 169 Nucleotides)

GITTIGCCAC ATTGGCCAGG CTGGTCTCGA ACTCCCGACC VVGTGAGCCA CCTGCCTTGG CCTCTCAAAG TGCTGGGATT ACAGGCGTGA GCACCACGCC CGACCCATAG CTCTTTACAA CTGCCTTGTA AAGAAAGCAT CATTTGGCAC TGTTAGTATT TCTCTTGAA

SEO ID NO:307: (Length of Sequence = 303 Nucleotides)

GATTIGGIAC AGAGIATGIC AGGAAGACAA CICAGATIGC CATITIAAAT AAAGIIGIAC ATGAACAATA ATIGGAATCA TCAGGIAATT TITITAAACA AAGGIICITC ATTIACIGIT ATGATTGGAA AAAAAATTAG AAAATAAAGI AAGISCCATA GGCTAATTAA AAAATAAAAC CIIGGCCGGG CGCGGIGGCT TACGCCTATA ATCCCAGCAC TITGGGAGGC CGAGACGGGC AGATCACGRG GICAGGAGAT TGAGACCATC CTGGCTAACA CGGTGAAACC CCATCIGIAC TIG

SEO ID NO:308: (Length of Sequence = 143 Nucleotides)

ATCTAGGAGG CTGAGGTGGG ATCGCCCCAG TACTGGAGGT CAGGGCTGCA GTCAGCCATG ATCATGCCAC TACACTCCAK
CCTGGGTGAC AGAGTGAGAC CCTCTSTCAA AAAACCTCAG TCAATVCAAA CATACAGTAT ATT

SEQ ID NO:309; (Length of Sequence = 199 Nucleotides)

CCCACCCTCA TAANCCCCAC TGGGGAGTCT GGGGGCCTCT ATTGCCATGT GCCTGGAAIN ATNATATGCT CATCACTTTA
TGAAGAATAA AATTTGINTT TCCTGCCTTA AAGTTACATT CGTTCTTCCG CTCAAATCCT GATCTGGTCC ATTAAAGAGT
GTTCGCAGAC AAAGTTTCTG AAAGATTAGA GAAGAATCC

SEQ ID NO:310: (Length of Sequence = 426 Nucleotides)

TCCCTGTACC ACCICTICCT GAATACGGAG GAAAAGITCG TTATGGACTG ATCCCTGAGG AATTCTTCCA GITTCTTTAT
CCTAAAACTG GIGTAACAGG ACCCTATGTA CTCGGAACTG GGCTTATCTT GTACGCTTTA TCCAAAGAAA TATATGTGAT
TAGCGCAGAG ACCTTCACTG CCCTATCAGT ACTAGGTGTA ATGGTCTATG GAATTAAAAA ATATGGTCCC TTTGTTGCAG
ACTTTGCTGA TAAACTCAAT GAGCAAAAAC TTGCCCAACT AGAAGAGGCG AAGAAGITCT TCCATCCAAC ACATCCAGAA

TGCAATTGGA TACGGAGAAG GTCACAACAG GCACTGGTTT CCAGGAAGCG CCATTTACCG TTTTTMATGG GMCAAAGGGA GTTACATTGG CTATGGCTTT TGGAAG

SEO ID NO:311: (Length of Sequence = 489 Nucleotides)

TOGACTOGGI COTGGATGIG GIGAGGAAGG AGTCAGAGAG CIGIGACTGI TITCAGGGCT TOCAGCIGAC COACTCICTG
GGGGGCGGCA CGGGGTCCGG GATGGGCACC CIGCTCATCA GCAAGATCCG GGAAGAGTAC CCAGACCGCA TCATGAACAC
CTTCAGCGIC ATGCCCTCAC CCAAGGTGIC AGACACGGIR GIGGAGCCCT ACAACGCCAC CCIMICGGIC CACCAGCTGG
TGGAAAACAC AGATGAAACC TACTGCATTG ACAACGAGGC CCTGTATGAC ATCTGCTTCC GCACCCTGAA GCTGACCACC
CCCACCTACG GGGACCTCAA CCACCTGGTG TCGGCCCACCA TGAGCGGGGI AACACCTGCT TGCGCTTYCC GGGCCAGCTG
AACGAGACCT GGCAAAGTGG CGGTTGACAT GGTGCCTTTC CTGGCTGAAT TTTTAATGCC CGGTTTGGGC CCTACCAGCC
GGGGAAGCA

SEO ID NO:313: (Length of Sequence = 302 Nucleotides)

CTICICATGC CAGICTAATG ATTGITTITA GAAAAGGATA TACATTGACC TTCAATGIAA TAAGAAATGC AACACTITAC GGTGICCAAC TGCTAAGATT TATTITCCAAC TIGTCAGACA CAACTATTIT GCCCAATCCA AATCAAAGGG AATCAAAGGCT GTGAAATCCA CACAGGACAT CAACGCACAC ATAAATGAAA ACTACAGATG TGTCAGAGGC AACCATATAC ACACAAATAA TGTAACTACT AAATTCCATG AAGTAGCTGT CCAGGGAATA CTTTCCAAAT AACCTTCAGC AG

SEO ID NO:315: (Length of Sequence = 339 Nucleotides)

CECETTATIT AAATTGIGAA AAATAATGAA TATTAATITG GAGCATAATA TITAAATACA TGAAAAAAGC TGGCTGGGAA ATGITGGCAT GACTTTCCC AGATGITAGC ACTGCTTCAA CITITGAGAG NGCACTCTGA GTGTAAGTIT ACTAGACTGA CATTACTAAA ATCATTGGTG CTATAGAGGC AGGAGAATAC GGGGAATAAG AAAGCCAGIT GCAAGCCAAC AATCCTAAAA CTCCTCCTTT TGCCATGGAC TGACGGCCATA TTAAATGAGA TCATGCATTT TAAGGATATA ACAGTGTACA CCACATGTGC GTGTTCCAAT AAAAGGAAG

SEO ID NO:316: (Length of Sequence = 430 Nucleotides) .

TAAGE: TIGGIG GIGCIGITCI GGATGCITCC AGIGGGCCCC GACCAGGICI GGACAATGCC TIGGCGCCCGT CCCCCGCCCC TCTTCTACAC ACACGCAAGA NITCGGAGCI CCATGGGGAA CAGAAGCAAG ATATCCGIAA AATCAAAGIC TAGGGGGIGG GAATGAAAAG GGAAAAGIGA GGAACGGGA GCCAAACCCA GGAAGACGCC TCTTTTCCTG CACATTCCCT CTCCTTTATA TACTCAGGCTC TTGGCTGTCT CCAGTATGTA CCCACCCTGG TCTTCCAAGC TGGGAGCCAC TTTTTATAAC ACAATCACAG TTTCACAAAC CCCAGGAAGG TTCCATGTGG NGAGAGGGTTA AGITTCGNCC TTGTCCGGGG AATTATGACA CTCAGAATAT CCCCTTTGGT GTAAATGGAA GACAACCTTT

SEO ID NO:317: (Length of Sequence = 317 Nucleotides)

GITAATGCIT CINATACCIA ACAAATCCIG GAGGGCAGNC AGCACCAACA CTCAGGGIGC TGGGAAAAGG TGCGTGAGAG ATCTGAGGCA TCTCGGGGGC AGGGGAGGGC TGGGAAGGCA GGCTGGCING GACCCTCGCA TCTTAACCIA ACCTTGACCC TCTTTCCATG AGCAGAGTIC CGATGCCCTG GAAGCCTGGG AGAGTGGGGA GAGATCCCGG AAAAGGAGAG CAGTGCTCAC CCAAAAACAG AAGAGTGAGG CTTCCAGGGT GCAGCAGGGG TGGGAGGTGA TCAAGCAGCG TGGGGATTGT AAGCCCG

SEO ID NO:318: (Length of Sequence = 407 Nucleotides)

CTCGCCCCGC ACCTTCCCCG CCTATGCCCC TCGCTGAGAT AGGCCCTTCC CTCCTCCGGG AGCCTCCCGG GCCACGCGAC
CCTCAACTTC ACAGCCGCT CCACCCACGC TTCCTGGACC GCCTCCTGCA GGCGAGGCTA ACATACAGCA CTGTCCCTTA

CAGTOGCCAT GCCCCTGGCG ACCTCAGTGT CCCACTCTGT AAGGGGACAA TGCAAATCCC TTTGCCTCAT AGGGTGCATG
TGCCAGTINTT GATAAAGTGC TGGCCACAGG CCCTGCCTTC CCAGGGCTCA CAACACTGTG TCCCTGACAC ACCCGTGGGC
TGTAGTGATT CINTTCATGG GGATTTGACT ATAACCNGCA GTCAGGAATG AATTTCACAN CATAGCTCAG TACATACACA
CATATCT

SEO ID NO:319: (Length of Sequence = 382 Nucleotides)

CACTGCACAC CIGCGGITGG GGACAGGACA TGACTAAGCA CAGAGCTITC TTCTTTTGAG GCCACGCATG TGGTGCAGAG
CGGGACCACC TGCATCCACA CAGCCCGGGG CACCTGCTCC TACTTCTGCT TAGCGTGTGA GCAGCTTGGT GACCAGGGTC
TCCACCAGGG GGCAGGCCAG GACCGGCTTA CAGCACTTTC TAGGGGTTCT CTGGTCCCGG GCTGGGACAC ATACAGGGCT
TAGTAAAGTT CATAGATGGT AGCTAGGCAG CCCCAGGCCC CAGGTGACAC CINTCCCCTG CCTGNCCTGT ACTGNCTGCC
TGCAGCACTC CTGGGAATCT TGTACGAAGA CAAGGAGAGA CAGGACTTCA TCTTCACCAT CT

SEO ID NO:320: (Length of Sequence = 368 Nucleotides)

CATCOGGGGC ATGGACAGCC CCCGGGGTGN CCGCCCGCNC CCCCCTCGCC GCGTCGCGTG CNGITCACCA GGCAGCACCT
GGACAGCTCC AGAGTCGGGG AAGCGCCATG GTTCCTGCGC AGAAAGGATG CGGGTTGGGG CCGGCAGATC CTGCCAGGAC
TAGGGGGCCTT CCCTTTCCAT CAGGAGCCTG CAAGAGAAAC AAGAAAACAT TAGAGGGGCT TCTGTGTAGG GGGAGGGCAA
GTTGAGTCTA TCTTTCCTCT TGTAGGTACT AATTAAACAC CTGCTGTNTG CCTGGTACTN TGCAGGGTGG GACAGGCATC
ATAGCAACTC ACAGTGGTCC CCTCTTCTTT GTGCCCCATAG TCTAGTAG

SEO ID NO:321: (Length of Sequence = 355 Nucleotides)

GGIGGACTGT GCTGTTGAAC TGAGCTGAAC TGGGATCAGG AGAAGGAGAA GTGGGGATTG AGCCCCTCAC CTCCACACAC
TCCTCTCTGT GCCTGAAATT CCTCCATTAA GCAGCATCGC TGTCCCCTGT AAACACCCAC ATTAAGCCAT TATTCATCTT
ATGGCTTNAG TAGGCGTTAG TCCCTCAGAT CCTTTCCTGC TGAAAGCGGA TCCTGATAGA GAGAAGGGAA GAGAAGATGGA
TGGNTCTGGG GACGGCAGGC TGGTCCAAGA GTGGGGAGGA AAGATGTCTC TCGGACTCTN GGGNAAGAAA TATTTTCTGG
GGGAATATGG AGGCACCANA GGCAAGCTCA AGAGG

SEO ID NO:322: (Length of Sequence = 225 Nucleotides)

CTCTCACTTC TCACCAGGCA CCCACAAAGC CCCCAGGCAG CTCCATCTTT CCAATCCANT CCCATTATCC CAATCTCTAC CCCAGGATCC CCCAAACTCC TCCCACTTCA CCTCTGCCAC AGACCCGCTC GCCCCCAAAC TTCAGCCTNC CCTCATCTGC CCTNACCACC CACAGCCCCT CCTACCTAGC CCTCTCCCGC GACGGGCCCG CGGGCTCCCC ACATT

SEO ID NO:323: (Length of Sequence = 250 Nucleotides)

CTCTCGCTCC TGTCCGTGAC CTTGCAGATG CAGGTGACAG CCTGCCCTTC CGTTTTTNTC TTTCCAGTCC CGCCTGCCGG
ATTGGGTTCC AGCCCTGCCC ACACGCCCGG TACATCCCGC CTACACTCAC CGATGTCGCC TAGCAACCCG GCTCGCCGCC
AGCATCCGCA ACCGAGGTCC CCGCGCTCCA GTTCTCTGGA GGGGAGGGAG AGGGGTGTTG CTTCTCCAGC CCCCTGCAGC
CTGGTGTCTT

SEO ID NO:324: (Length of Sequence = 338 Nucleotides)

GIMTCTTAT GCGGATAAAA TITCINAGGI AAGAAAAGIT AGCTCTGAGC AGCCCTCCGC CTGATACTAA TACTTTACCA ATGGAGATIT TYCTTTTCTT TICIGITTTT GAGACAGGGI CTCACTTGI TTCCCAGGCT GGAGGGAGI GGGCCATCA TGGATCACTG CAGCCTCCAT TTCCCTGGCT CAAGCCATCC TCCCACCTCA GCCTCCCGAG TAGCTGGGAC TACAAGGTGI GCACCACCAC GACTGGCTAA TTTTTAATTT TTTNNTAGAG ACGGGGGTTT CCCTATGTTG CCCAGGCTGG CTTGAATTCC TGGGCTTCAA GTGATCCT

SEO ID NO:325: (Length of Sequence = 461 Nucleotides)

SEO ID NO:326: (Length of Sequence = 391 Nucleotides)

GGCCCTCCAG TGTCCTGCAG AGAGGCACTC TTGCCAAGTG TCATTGATGA CGCAGCTGAA AACCAGAAAC ATTTCACTTT
CCAGCCACGA GACTGCAGCA ATCTGCTCTT TGGACTGCAC TTAGGGAAAC CGAGGCCCAG ATAACTGACC CCTCAAAAGC
CCCCAGGACG GCAAAATCAA AGGGGCTGAG GTGCTCTGAA CAGCCCCAGC AAATTAAACC ACCTAACTTT GCGCTACTCC
CACTGCCCTG AAGCAGCCTG TGGTGGGAGG TGGGGGTGAA TACAGTGTTA CAAAGAGAAA CCTGAGTTGT AGCCATAGAT
TGCTAATCAG TAACAAAATA TCCCTCTAAA CCCAGTCCTG CCTTGAACCC ACAGGCTCAG GATGGTAAAT A

SEO ID NO:327: (Lementh of Sequence = 438 Nucleotides)

TACTGACTGA CCCTGG: GATTCCCAGC CGAGACGTTT CTGCTCCATT CCGGCAGGAG CTACCTTCCC GAGCCGCGCT
TTGCTCACCT GIAGGAG: TAGAGGGAAA TAAGACAGCC CTTCTTAGGA TGGTGGAGTG GCTAGAAAGA AGCAATCCAC
GCCAAAGGCT TAGCTCAGTT CCTAGACTTA GTAAATGCTC AATAAATGTC TGCCATTGTT ATTATTATTT ATNATGCTTC
CAGCTGGCCT GGAAGGAGGG TTCTGGAGCC AGAAGGGACC TTGGAGAGAC CTCGGTTAAA TCTCTAGCGC CATCTTTATT
TTTAGGATGG AGTAACTTGC TCAGGACCTA CATCTAACAT TGTGGAGGGG ATGCGGTTTT TAAGTAGGAA TTCTTNGACT
AGACCTCTCA GCAACCCTTT CCTNTCCGTG ACAGTGGG

SEO ID NO:328: (Length of Sequence = 400 Nucleotides)

TIGCCCTCTC GGCCTAGAAG TCTCCCATTA TGGTGCTGTG TCTGCTGGGA CCCACGGGGC GCTGCACAGG GAACCATGTG
GCCGTGAACC TCAAGTCCNG NCCAGCAGGG GTCAATTGTC TCAGNCCACC CCTCCCTACC CCCAGTATCC TCTCTCCTTT
ATAGATCATC CATTAAGTGC CAGACACTGC AGAAGGCACA TTGACTAATA TTAAATATTA GCCCAGCTAC CCTGCTGGGC
TGTCCTTCTT AGAAATGAGG AAGTGGAGGG TTAAGTGGAT TTCTCAAGGT CGTGCAGCTG GTAAATGGCA GAACCAGGAT
TTGAACTCAG GTGTGCATGA CTTCAAAGGA AGACACCACT GAGGCCTCCT CTANTGGGTC TGCNTCCCTA CCGGCCCTGG

SEO ID NO:329: (Length of Sequence = 227 Nucleotides)

GECTIGGECTA AACTICCAGAC GCTGGCCACC TTCATAGGGT GEAGATGACA GAACAGGACA GGAGCCATGG GGCTCCCGGG GCGGGTAGGG GTGGGTCATG TTCCTTGGCT TGGGGGCCAGT TACAAGGGTA CAGTGGGGCT TGTTGAAGGG CAAAAGTTCT GTAAGTNCGT CCCNACAGGC CAAAGAAACC CCAGAGCCGT CTTTCGACTG ACTACAGCCT GGAAGAG

SEO ID NO:330: (Length of Sequence = 401 Nucleotides)

TGAAAATATA TOCACTIGTIC AGAGGGACAA CAAAGGCAGT TAGACTIGTOC TGAACEGTOC TGCCTCAGGC TGAAATTITT GTAGCACTIG ATCAGTIGCA AAGIGATCIT COCTITAATA TCTCATTITA TCATTGGGTA TCTGAAGAG AAGIGGAATT GGGGTAAGAA TITAGGTTCT TGCCATAGCA TITGGGTSSC CAGGGTTAGC CTCACGGTG AGGACCCTTA AAGAAAAACTC TAAGGATTIT AAGGAGAGIC AAACICIACA TICATCCAGG CAAACATCIA CICITCCAIT GATIAATGGN TCCACICATC
CGIGCAACAC AITCACICIT TCATCCATCC ATTCATCCAT CIATCCINCA TCAATCCATC CATGIATCIT TCATTCATCC
A

SEO ID NO:331: (Length of Sequence = 322 Nucleotides)

COCAACGITG CCCCGCCTT GICTCCAGCG GACTGGAAAG AACCCACCAT TGTGAAGCAC AGAAAATTGC CCGCACTCTT
ATTGGCTAGG TTCCCCGACT TCCGCTCTCG GITGGTGGTT GGCTTTGCCT GITACCTGTG TTGCCCACTA CCACTCGCTC
CGCCGAGCCC CAAGGATGGA TCGCTATCCC GIAGCCGGGT GITCCGGAGC GCTGCGGGCA AAGCAGACCG CCTTGCGCCT
ATTATGGGTT GAGTGGCTCT GIACTCTAGA TCGGCTCTGT CACTTACTAA TGGGCCGTGT TGCCTTCGCG ACTGCAGGTT
TT

SEO ID NO:332: (Length of Sequence = 441 Nucleotides)

GECTCAAGNA ACCTGCACTC TTGCACTCTG GCCTTCTCCC AGGCTGAGCT TTATCATATC ATCAGCAGCA ACCTGGAGAA
AATTGTCAAC CCAAAGGGTG AAGAAAAGCC ATCTATGTAC TGAACCCGGG ACTAGAAGGA AAATAAATGA TCTATATGTT
GTGTGGGATTC CCTTCTGGGG TGTGTCATTC ATTCAAAAAG CATTTATTGA GTGGCACCTA TGTCCAGCCT GAAGATGAAT
GTGGTGGGAA GGGGTGGGTG TCACAAAGAC AAAGATGACT TAGATGCCCA CTGTAATCTT GACTGTGAGA AAGAGGGGAT
TCAGGGCCCTT TCTCATCCAG TAGTCAATGT GCCATCTCCC CTTCCCTAGT CACCTCTTAT CTTCACTTAC CTTCTTTCTT
CTCCTGCTTA TCTGTTTTCC ATCTAAGGCA AAAAGGGGGGG

SEO ID NO:333: (Length of Sequence = 354 Nucleotides)

AGRAGOGTAG ACCEAGTAGC TTGAGOGCCT CTTCCGGTTA CCTTTTCCCA GOGCCAGAGG GCCTTAGGGT TGGGGTCCTC
GCTCAGGCAC AGAGNOCCGA CACCGAGCGG CGGCTTCCCC GGGATCGAGG GACGCGCACG CCAGAGGAGA CGAAAGGAAC
CCGGGTCGGA CCAGATCGGA ACCACTGACC ATTGCCCATG GCGCCCCTAG TGAGTNTGGA TTTNGCGGGG TTCGGGGGTT
CCGACGGCGA CCTCGGCGAC CCCTCACTCA CCGCTTCCTC TTTNCNCAGG GNCCTAGNAG CCAGAATGTC ACTGAATACG
TMGTTCGAGT TCCTAAGTAA GTCCCCAGGC CCAT

SEQ ID NO:334: (Length of Sequence = 196 Nucleotides)

CTCCCGCTCC GCACCCGCCT TTCCCGAGCA GGCTACACCT CTCCCTGGCG CATCTTTACT GGAAAGCCGG CAGNGGNGNG GGAGAAGTGA GCNCCGTCTC CGCGCCTCCT CGGTCCTGCT GGCTGAGCGC GGGGATGGCT CCGGAGGGAG ACACTCAGGA AACCACCTCC GCCCTTCCCC CATCTTTATC CAGCGG

SEQ ID NO:335: (Length of Sequence = 261 Nucleotides)

TOCGAGAGCT GICTGGGGCC AACGIGCTGG CTGAGTACTA CTGGCTCANA CGCCGCCTGC TGGGGGCCCC TGGAAATNTA
AGICCTGCCC CGGGCTGTGC CGCCCTCCTC CCTGANAGCC CCCTGCNTCC TGGGCACAGG GAAGCCTCCA TAGGCTAGTA
GCATCACAGT GCCAGGCCCA GAGCTTACTG GACTTCCCAA GGTCCTATGG GACTAGGGCT GAGGGTACAC ATCCTGCTTT
TTTCCAGAAT ATAAGTTTTG G

SEQ ID NO:336: (Length of Sequence = 191 Nucleotides)

OGGAAAAGOG CTTOGGCCAC ATCCAGCAGC AGTAGCAGCC GCAAGGNCOG GGACTGGAAG GCCCACCGNA GNOGGACTAA GTOGTCCAAG GAGCOGCCTT CGGCCTACAA GGAACOGNCC 32.252777ACC GGGAGGACAA GACOGAGCCT AAGGCCTACA GGCGGCGGCG GTCCNTCAGC CCACTGGGAG G SEO ID NO:337: (Length of Sequence = 279 Nucleotides)

CCTTAGGGCT CCTCCTGACT CCTTCCAACT CCCAAGTCTG CAGCCCAGGT AAAGCCAGAG CAGACTNAAG GCAAGTTTTC AGGAAACCAG GNGGCTTGAT CCAGACTCAC AATCTCCCTG CAAAAGTKTT CAGAACACAC CGCACAAACA CACACACGNC TCACAAAACT TCTGAATGTK GCTCTGTCTC CACCTTCTCC AGTCACCGAA AGACCTCGGC CTGAATTGGA GCCCGCAGCC GTAGCTGTCC CINTCCACCT GTNGCCCTCG CGGAGGCTT

SEQ ID NO:338: (Length of Sequence = 339 Nucleotides)

CCACNOGIGE AGGRAGICA AGGGGCAGCA AGAGAGAGGG AGGAAGCCCC ACTITITIAA AACAACCAGA TOTOTIGIRA
ACTGAGAACT CCCTTATCAC CAAGGGGACG GIGCTAGACC ATTCATGAGG GWICCGCCTC CATGGGCCAA TOCCCTCCCA
CCAGGCCCAC CTCCAACACT GGAAATAACC TCCCAGCAGG CCCGCCTCCA GCACTGGAAA TAATGCTTCA GCGTGAGACT
GGAAGGGGAC TGATGGAGCC TGGWTGTTTK TCCCCGCCCA GSTCTMACGC TGAACCGTAA TCCCCAATGC TGGAGGCGGG
GCCTGGTGGG AGGTGACTG

SEO ID NO:339: (Length of Sequence = 334 Nucleotides)

GGCACCGGGC TGTCTCTNET CCAGCTAGCC TCACAGGGAG TGGCCTCTAA AACNGGCCGG CCCACNCCAT TTGGAAGCTG
TCCCGGGTTT TCCGTGAAGT CCTCCCGGCC TGTGGTCTCC TGGATGGTCT GGACCAACAG CTTGGGGATG AGGGGAGGCT
CGGGGGCAAG GGCAGAGCC CCAGCCAGGC GCTGGGGGTN TGGCTGATCG AAGAGCTGCA CCACCCNGTA GCTGGCCAGG
TGAGTATNGG CGTCCACCAG GTGCAGACAC ACATTCTTTT CCTTNACAGC CTCCTTACCC TGGAGTTTAT AGCCAAACGT
GAGGTCGATC CAAT

SEO ID NO:340: (Length of Sequence = 450 Nucleotides)

GECCCCACAA TCCCCITCIG GCTCCGGGGA CGGGGGGGAA GGGGGGAAATA ATTINITGIT TGGTCGTCTC
TGCCCCAGTC CCTTCGCCGC GGGACGGCGA GACGGGAGAA GGTGCGGGAA GCGGGAAGCA GGAGCGGGAG CGCCCGGCCC
TGGCACGCAT AGGGCGGCGG AGAGGGCACG AGCAGGGATT GAGCACCTAC TGINIGCCTT CACGCTTTAC AAAAGGATTT
TCGTTCGATG TTCACTACAG CCCCTGCCCG GGGGTACTGA TGCCCCATTT ACAGAGGGAC AAGCCGGATT TCGGAGAGGT
GAAGTCACTC GCCGAAAGTC GCACCGCCAG GGTCTGCGTG ACACCCTAAA GCAGTGTTCA GTTACCCCGG GGAGAGCGCG
ATGAACTTGA ACCACTTGTT GGCTTGGTTC CTGCTCTTGC TCGTTTTTTT

SEO ID NO:341: (Length of Sequence = 192 Nucleotides)

TTCAAACCCT GGCGGCACGG CTTGTCCCTC GAGGCCCGGC CCCTTCCCCT TCCGGAGAGC CCACCGCTGG GTCCTAAAGC CCACCGCTGG GTCCTAAAGC CCACCGCTGG GTCCTAAAGC CCACCGCTGG GGAAACCAGG TCTTTCCTAG CTCTTGGNTT ACTTCCTGGA GACTTCTTAA AACGAGAGGA GA

SEO ID NO:342: (Length of Sequence = 229 Nucleotides)

GIGGIAACTI TITTAAAAAA CATAAATACC ATACAATTCA TCCITTTAAA GIGIGIAATT CAGIGGITTI TGGIATATTC
AGIGITGCAC AGICATCACC ACIAATTCCA GAATATTTIC ATCACNCCCA CGGCTGTATC TCCCCATTTCT CTCTTCCCKG
CAGATCCTGG CAACCGCTGA TCTACTTTCT GTCTCTTACA GACTTATCTG TTCTGGACAT TTCACATAA

SEO ID NO:343: (Length of Sequence = 229 Nucleotides)

TECTOCAGGA AATTGGAGIT CNAGCTGAAG GCCTTGCNGC ACTCCGNGCA CTCGTAGGGC TTCTNGCCCCG TNTGCGTGCG
TCGGTGCTGC ACCAGCGTGG TGCTTCGNCC GAAGACTTGC CGCAGTCCCG GCAGCCGAAG GCCTTCAGGC CGCTGTGTGT
CTCCTGGTGG TGGATGAGCT GCGAGTNCGC GCGGAAGGCC TTNCCGCACT NCCTGCAAGC GTAGGGCTT

SEO ID NO:344: (Length of Sequence = 227 Nucleotides)

TCCGCAGATC ANATTCACCC TTGCCAGAGG TCAGGSCCCC CGGCCTTGGC GGCGGGCCAG AAGCGTGACT TGGCCTSCTG
GAATGCATGC CCCTAAACAT CTCTAGACTA GGGGCAGTKT CCGCCAACCA TGGAGGCCCT CCATCACCAT CCCTGCAGCA
TCACCACCNT CCAACCCCCA TGTCCCACCC TGGNGNTTCC ATACCTGTAG TAAGAGAGCA AACCATT

SEO ID NO:345: (Length of Sequence = 249 Nucleotides)

GGGCAATGIT GICACAGATG TGTGCAGATT TTSCAGAGGA CATAAGITGG CIGTGAGGWA GAACACAGAG GITSCCTATT
TTTTAGGCAG GAAAGAAAGC CIGCACTITT CIGTGTGTGT GINICAATAA ATCIGAATAA CACCITGAAA GGGTTAAAAA
GCTGAGCACC AGGTGTTTC TTTCCACTIT CCAGAGTAAT TTAAGCACAC NSCAAAGITA TCTCCCTTCC TTCCCCACGA
GCCAGCTTA

SEO ID NO:346: (Length of Sequence = 356 Nucleotides)

ACCTAGICCC GCAGCOGCTG CAGCOGCTGG GTTGGCGGAA GAGCTGGACG CCGAGCTAGA GGACGAGGCA GAGCTGGACA
CAGTGGCGGC GTGAATTGGC CACTINCITTC GGAGCCCGAN CTCTCCCGCA CTGGAGAGGA CTTCTTCTTG GCTGGGCGGC
TCTTGGTTCC GCTCCCGCTC TGCTGCTGCT GGCGGCATTT NGCGCGGCGG TTCTTGAACC AGACCTGCAG TGGGCCGGAT
GGGGGAGAGT GGGTCAAAGG GAGCTAGGGG AGCTINITGC TCCACCGNCC CGTGGACCCA ACTCCCGGTC CAGAATATCG
CAATCCTTTC TCACCGAGGC CTTCGACCCT TCCTGT

SEO ID NO:347: (Length of Sequence = 155 Nucleotides)

GCCGCGGTGC GICGGATGCC CAGCTCGCGT CCAGACCCGC GGGATGCAGA CCCGGTTCAG TCAGGCTTGA GGGCTGCTCC GCATAGACCA ACGTCCGGGG AAGGCACACA GTGGCCGAGG GCCCGCGGCC TTKGGCTACG GCTGTRATGG TATCT

SEO ID NO:348: (Length of Sequence = 362 Nucleotides)

AATTOCGATT TAACTGATTG TCTCATTCTG CTCATACATT TCAAGTTTAA ATGCAAGCAT AAAATGTTTA TCAACAAATC
TAGAGAGCAC TIGGATTTIN AATTTTCCTG TGATCACAGT AAGGAGCATA AAAAAGAGTA TCINCTGTTA CACAAGGCCT
GINCTCTCTT TACATCTTCA GACTTAAATT CTGTAGAAGG TAACAGCTTT GTATTAAGGA CAGAAGCTTA GTGGTCACAA
ACAAAAAATA ACACTGAAAT ACAATTCGGG NATTANTGAT ACTGTGTGTC TCAAAGGATA CCTGAACTAT TACANTNACT
AATAATTTGG GCAATGAGAT TCCCNGGTGN TTCAACTTTT TG

SEO ID NO:349: (Length of Sequence = 342 Nucleotides)

AATTCCTTT TITTTTTTT TITTTTTTT TITTTTTTT TITCAAGTAT CACAATGITT ATGATAGAT ACAAGTATAT
AAAATCAGGG CATGANCATG ACTTGATAAA TTAAGTAGAC TTAATTTCAA TACTATAATA GGNGGGACCA ATTCAAATTC
TCACCATTTG TITCACACCC ACAAAAACCA CTTCAAGGGC ATTAACGNTC TCTCAAAACT GNTCAGTTTT GIGCAAGTAA
ACCATGTTTC TITTTAAAAAG ACTTGTGCAC TTGCCCAGGC TCAAGGTTAT TAAAATCTAG GCACATAAAG NCCATTACTA
GAGGTAGGAA ATACAGGCAA TT

SEO ID NO:350: (Length of Sequence = 384 Nucleotides)

GATCTGTGCT AGCTGTGAGG CAGCTCTGGA ACGTGAAGAG CTGTTTGGTT TGANCCGTGA ACAAAACTGT GTTTTGAGTT
TAGCTGACAT TAAAGAAAAA AGTTCATCAC GTGACTGTTA ATGTAAACCT GGTTATTAAA ATAACTATTT AAAACAGGAG
AAATCTGGTA AGTTGTTAGG NITCTAAATT CCTTTTAGTC TGTTCACTGA GATATTAAAT TTCAGTAGAC AGAACCCAAA
AAGAGATTTC ATTTCTTTCT AATCACTTTG GCTTCTNTCT NITTTNTTAA GTAGGTAAAA ACCTTCCTTG GTGGGCACCT
AAGCAGGATG CAGCCAATTA GTTCATGAAC CCAGCTGCGG ACGTGAAGGC TTAAAATCTA AGGA

SEO ID NO:351: (Length of Sequence = 305 Nucleotides)

ATCCTGACCC TCCCCACTGC AAGCCCAGGG AGCCCCAGCC CAAGATGGCC AGCCTGAAAC TGTTGGCCAG GGCTCCTCTT
GTGGCCATGT ACCCAGGGCT GGCTGGCCTG CCATTTGCCT CTCCCCGGAG ACAGCCGTTC TTCTGCAACC ACACCCCGTG
CCTAGCCACA ACCCCAGGCT GCAGCTGCTC AGAAGCTCCA GGCATTTTGT TTCTGGTGAC CGCCCCTAAT GGGATATCGG
TGATCACTGG TCCACCCTTC CTGTCAGGGC TTTTCTGGGG GCTGCTCTTG GAAATGAAGT CTTAA

SEO ID NO:352: (Length of Sequence = 270 Nucleotides)

GAAATTACCC ATGGTCATAT CTAGCCTACA AAGAAGAGAA AATACAGTGA TTCAAGTTTC ATTGTATTCC TCTCATTGAT
ATATTTATCA ACCTTCCAAT TGAAGGAAGT GTCTTCTAGG CCTTTACAAA GAATGTAACC AGGGTTTAGG TATACAAGTT
GCATATGATA AATCTGTCAT GTTTCTATAT AAATCTGTCC ATATTCCTCT TCTGAAATGC ATTATTTTTTG GGGGAAATTA
AAATGTGATG CAAAGATCCT TATACTTTGT

SEO ID NO:353: (Length of Sequence = 195 Nucleotides)

GIGIGATICC ATTIATATGA AATGNCCAGA ACAGGGAAAA CCITATTINAG ACAACAGAGA CACAAAGTCG ATCAGCAGTT GCCAGGGGAG GAGGAAGACG GGAGGGGAAA TNATIGCTIC ACGGGGTGAT GACAGAATGT NCCAGAACGI GACAGAGGTG GIGCCTACAC AACTITNIGG NIGIACTAAA TGCCG

SEO ID NO:354: (Length of Sequence = 388 Nucleotides)

GCCAATTITT TTATTTTGT AGAGATGGAG TCTCCCAATG TIGCCCAGGC TGGTCTTAAA CTCCTAGGCT CAAGGGATCC TCCCAGCTGG GCCTCCCAAA GTGCTGGGAT GATAGGCATG AACCACCATT CCCAGCCCAT TTCCTTTTTC CCTTTGCACA GTACCAGATA TATGGTTGGT ACTGCAGAAA TAATTTCCCC CTGCCCTCTA CATTGATCAT TTGATGACCA AATAGTGTCC GTCTAGCCAC TTATTTATGA TTTGTACAAA ACATTCCGCT TTCTGAGGTA GACAGTGATA TTCTGAAGCC ATCAGTAAGA GTAATTTTTC AGINTTGTG AAAGTGCACA TTCTTTGTGT AAAGGTCAGC CTGTCAAGGA AATAGCAT

SEO ID NO:355: (Length of Sequence = 288 Nucleotides)

TAAAGIGAAG TATTGGGAAA GGGAACATCT CACTCTGATA GATTIGAATT INCTATTTCT GCTCTGTGAC AAAACCCTGA
GTTGATATGT GATCAGACAT TTACAAGGCC CTGCATTCTA CCTGGAAATG GCTATAGIGG TGTTGAGCTG CTGTGAGATG
ATTTACTGCA ATTTGTCACT TTTTGAAACT GTTCCAAAAT AGTCTGCTGA CAGCCCTTCC CCTCATGAAA ACATCTCTCC
TTTTCCAGTT AAAAAAACAG TCAAAAAACA CCAAAAAAAGG CCACCTCC

<u>SEO ID NO:356:</u> (Length of Sequence = 401 Nucleotides)

GGAAATTAGG TTGGTTATTA ACATGIATAG ATGGAACTGG GGTGAAAAAA AAAAGGAAAT GGGAATGGAG TGGAAGGGTT
GGGTGGGAGA GACACTTCAC AGTATTCTIT TTGTTTTGAC TTTGGAAATG TTACTATTTC ATAAACTTAA AAAAATGCAA
AAAAAAAATA TCAAAACTAG GTAGGAAGGA GAACAAAATG AAATATAACC AGAAAGGAAT AANCCTAACA CATTTTGAGT
GAATCACAAA GCCAAACCAA AAAAGAGCTA ATTTAAGTCA CTTTTAAACT TGGTGTTTAA CTACCTACAC TCAGTCTAAA
AACGGNAAAT AAGGGTAAAG AAATAGTGGA ACTCTAGTTA GTTGGGTCTT TTCTTTACAG CAGTATGGGG ATGGCAACCT
G

SEO ID NO:357: (Length of Sequence = 275 Nucleotides)

CAGACAGTGG ATAATAAACA CCTCATTAGG AAACCGATCT CAGAATGANC TCTGGAGTAT GAAAAAGATC ATTTCTTTTT
GINCCTGTAA CCTAGCATTC CTTCTAGGCT TCINCTCCTT TAATTGAACC ACAGCTTAGC TCATGTATTC TTTTATTAAC
ACCCTGCTCT CATGTCCATA AGATTCAGGA ATTTAGGAAA TNAGGCTGGT TTGAAGACGG TAGAAAGCAA TAAAGGCACN
AAAAAATAAG NCTAAAATCA GGGGAAGATG TATTT

SEO ID NO:358: (Length of Sequence = 314 Mucleotides)

GIGAAGGAAG TATGAAAACT GAGACIAATA TIATGAAGIC TITTTITAAT TCITTATCIT ATIGCCCATT TITAACCCCT
TGGIGIITGA AATGGAAAAT AAATAINCTC TICGCGATAG ATAATATGIC AATAACCAAA AGGIGGCCTT AACCAATAAT
TGGCCCCAACT TIAAATTATT ACCCTAAAGA TATATAAATT ANCTAATCIA AAATTAAATG CAATTITGCT ATGACTTAAA
GIGICANIAA TCCIGIATAA GAGATCCNNI TIATGCAGIC ACTTAGGCAT GAAGITGGCA ATTCATCTAA ACTT

SEO ID NO:359: (Length of Sequence = 372 Nucleotides)

CAAGAGAGAC ATAGCAGGCA TIGAAACAAT GGAAATGCCC ACATAGCAGA AGGGAGTGAG GGGATCCAAA CTACAAGAGC
GACAAAATCA ACTGTGGATC CAGAGACGAA AAAATGTTCT GTAGTGCAAA GGTAATCTGG TGAGATGAAA AAAAAAGAAC
CATTTTTAGA AAAANGGAAT ATTAGAAAATA TTGAAGTAAA TATCATAAGT CATTCTATTA CAAAGGCATT AACTCCTTCC
TATCAATAGA ATGTACCAGT TTAAAANTTT TTAGTAGGAA TATATCTTTT ATTTTATTAA CAGAAATCAN GGGACAAAGA
GGATTTGATC CATCCATACT TCCTACTCTT ATTGGGTTTG TCAAAAATGTA GG

<u>SEO ID NO:360:</u> (Length of Sequence = 395 Nucleotides)

GCATTCTITT GATACCCACC TAATAAAGAC AATCTCTAAA ACCAAATAAT AGGCTATGAA ATGTATTGTG AGINCITATT
TCATTCAAGA CAGAGCTTAC CTTTAAGTCT CCAGCTGAGA CAGTTGGTTT TATCTTTCTG AAAGCAGTTT GGTCAAGTGT
TTCAAGTAAA TCAAAAGATC GGTTAATCAA TTCCTTAGCG AATTGGATTA GACACTCTCA TTTCAAATGG CAGTTTTATG
CTTACTCATT GTCTTGAATA ANCTTAAATA CTTTATGCTA TCTTCCTGCT CCATTATTTA TGTAATCACT GGGNCCTTAG
TATTCTGCTT TAGNNCATAT AAAATCACTT NCAGGTATTT TCCATCACGG ACACAGAGGC AGGCACAAAT TAACC

SEO ID NO:361: (Length of Sequence = 298 Nucleotides)

ATTITITIGI GGGGAGAACA TITAAGACCA TITCAATGIC ATGATGAAAG CTAATGGGAG AAGGCITTIN INCIACAAAA AITINCITIA TITTINCAAC TITAITGAGG TIATAATTGA TATTAAAAAA CIGIACAGAT TIAATGIGIA CAGICIAATG AGITGGGACA TATGCTTACA CCCNIGATGC TGITACCACA GGCAAGGTAA TACACATATC CGICACCTGC AAGAGTITCT GIGITTCCCN NIGITTCTCA TITTGNITTT TICAAAAATT TACITTATAG GCITATAG

SEO ID NO:362: (Length of Sequence = 437 Nucleotides)

ATGCTGGAAG TGATTTCTGC AGCTCAGGAT TTTTTTTTTA AGCTACATTG AAAATATAGG TTTATTTTTT GTNCAGGTTT
TNCTTTTATA TTTTTTNCT GCACAAAGGA GGAGGATTTT CCACTTACTC ATATCGAGGC CAGATTTTTA AAGCCAGCTA
AGGCAGCATC AGCTGGCGG GATTTAAAGC CTATAGCTCA GCTGAAAAAA AAGGTGGGGT GGCGTTTCAT GTAATGGGAC
ACGATGCCCT TCTTGCTGAA CGACTGGAAA GAGCACAAGG AGCACTTTTC CTTCTCCACT GCCCGCCGGA GTTCCTCGCT
CAGCTGAGGG GAGTCGTCCT TGGGCGGGGA TGGGATGATC ACTTTGTTGG GCTTNTCGCT GATGGTCCTG GAGGCTGCCA
AGAAGTTGAG GTGTAATACG CATCAATGTC CGTGGCG

SEO ID NO:363: (Length of Sequence = 449 Nucleotides)

TGATTIGAAG TAAGCTITCC ATGCTTCACT TAGGGTGGGA AATTITAAAT ATCAGAGCTI TCTTTGTTAG CAGCATATAG
TTATGCAATT TATTTAAATC TGCAGTGCCA ATCTTTTTTT GATGGGTGTG CTTAGACCAC ACATTTAAGA TAATTATTAA
TATGTTAGAA CCGAATATAT TTTNATGATT AGTTTTTATG TGTCAATTTG ACTGAATTAA GAGATGCCCA GACAGGTGGT
TAAAACATTA TTNCTGGGTA TGTTTGTGAG GATGTTTCCA GAAAAGGCTA GCATTTGANT CAGCAGACTG AGTAAAGAAG
ATAAAGATAA TACTTGTCAT GTGTACAGGC ATCATCCAAT CTGCTCAGGA CCCCAATAGA ACAAAAAGGT GGAGGGAGAG
TGAATTATGT CTACCCCCTT GAGCTTGGGA CAGCCATCTT TTCATGCCC

SEQ ID NO:364: (Length of Sequence = 282 Nucleotides)

GACTGTGTAA ATACACTTTA TTTTCCATTT INCCCGCCTG GGCGACATGT GAACAGGCAG TGTGCAAAAT GGTGGCGGGC
AGTGTAGGGG GCGTGTGGAG AGCCCCGTGG GTENCTGCCC CGGTCCCCAG GCTTCGTAAC ACTGAAAAGT GGGCAGCTAG
GAAGCGGGGA CGGAGCAGGG GTCCCCACCC AGGAAGCGCC AGGNAGATTN CTTGTAACGC TACTCTACTG GAGGCTCCGG
GAGCACCGAG NGGGGCAGTC CCCCAGGGTCA TGAGGCCCGG GG

SEO ID NO:365: (Length of Sequence = 349 Nucleotides)

TTCAAGCATT TCTCCTGCCT CAGCCTCCCA AGTAGCTGGG ATTTCAGCAC CTGCCACCAC GCCCAGCTGA TTTTTGTATT
TTNAGTCAAG ATGAGATTIT TGCCATGTTG GCCGGGCTGG TCTTGAACTC CTGACCTCAA ATGATCCGCC TGCCTCAGCC
TCCTAAAAGTG CTGGGATTAT AGGCATGAGC CACCACACACT GGNCTTTTTN TTCTGTTTCT AACTGTTCCC TTTTATTTCC
CTATGGAGCA TCTACTGAGC CCCAGCCGAG AGTAGAAACA AACCTGCTGG CTGCTCTNAA GGCACTTATA GTCCCAGTTA
GGGGNNGACG GGTCACTTAA CCACTTAGT

SEO ID NO:366: (Length of Sequence = 366 Nucleotides)

ATGCAAAGGA ACAATGGTGT TGGCAAAGTC TTCTTTGAAT ATCAGAGACT GAGTCAATAA AAAAAATAGT AGAAAGGTGG CTTTTACTAT TGACAAAAGC CGGGTCAAA AAAAGTAGTT TAAGTCTTAA GNCTGAATAT GCATTAAAGT ATGCAGGTAG CAAAGATGTA ATAAATTTCC TTAAAAAAAAG AAATTAAAGT TTTATTTAGA ATCAATTTTA CCNGTCATTG TAATTGACCC MTCTGAGNAT TACAATAAGC AAGAGGAAAT TAAGGTGTTT TGCAAGAGCT GTATTTATAT TACNGNTTT TAAAAAACCAT TTTCTGAATT ATCGTAATTA AAGCTCTCCC AACTCGTTTA AGTCAG

SEQ ID NO:367: (Length of Sequence = 391 Nucleotides)

GCAAAAACAA ACAAACAAAC CITTAAGTAC AGTAGITCCA AAACACACTG CTAAAGTTAT GAAATAATTG TGGATCATTT CAAGTAAAAA TTATTAAAGG AGCAATAATT AACCACAAGG GGGCATATAT ATATTATNCNC CITAGATTCC AGCAGAAAGA CTAGITTTAA GTAGITAACAT GCACGITGAA GTATTCTACA TTTTCAGTCA CITAAACTIT CCTCTCCAG ATGGCTACAA CITTTTAATA TTCGAGGTNT ATTTTATATC TAAGTAAAAG GATTCCAGAA TACTCCTGCC CTGCAAAACA GTAGTGTTTT

AGAAGNCICT NGGAAGIGIT GCIGITTACC CTTTAGCAAA GNGNTACAAG AGCTATTAGT TGTAATAATA C

SEQ ID NO:368: (Length of Sequence = 370 Nucleotides)

ATTICCCITC TECACTOGGT TCTCCTGCTC CCCATTTACA TEGITTACTT CATTITCCTC TICATCCATT GGATTCACAT
GIGITCIAGG CCAATATTCC AGGNGTGCTT GGAGTAAAAG TCCTCCTAAA TTCAATTTTG GNICTGACCC ATCAGGGCTG
CTGAAACCAG CATCTTTTGC AGAAACCCAG GCAGCAAAAC AATCACTTTC ATCCAAAGTA ATAGTTAACA TCCCTGTTTT
TAAGTCTACT GAGAACCAAT TTGGCACATA CACCATTTTA AATCTTTNCT TAATTTCATC TTCAAAATCC ACTTTGCCCA
GATCTTCAAC TTTACATGGC TTCAATACAT CCCAATATGN CACCATTATTA

SEQ ID NO:369: (Length of Sequence = 315 Nucleotides)

GACAGGIATT CITTAGAAGI TITTIGITIA CITATGITTI NCICITTIAC ATCICCITGI GAATITCIGI CCCATTITGA
AGICTCICCI TGITCICGAC CAAGAICCCC TIGATGITCI GIAGCCAAGA ACTGAGAAAA AGAGITATIC TGAATGAIGI
AGAGGITGAT AAGICIGGIA AGAAACIGIT GGACATACIC CAAGCAGCAC TGCATTGCAG TCITTIGGGC TGICTICCTA
CITCGGGITG CIGICCCCIG AGIGACIACG GAAGGGGICT GGATGATGGI TICTICAGAT CCCACAGIGG ATGCT

SEO ID NO:370: (Length of Sequence = 442 Nucleotides)

AACACTITIA CACTGCTGGC CTAATTIGIA GATATCCTCA AGAAGATTAT GAGTCATTCT CACTACCGGA ATCTGTTCCT
CTATTTINIT TACCAATGGG TGCACCATTG AATGTTGGCC ATCAAATAGC AAATACCCTC TGCCTGTATT TCCTACININ
GTTTTAACTG GAGCCTCAGC TGAAAAAGGIT TATGGTGCTG CTATTCAGIT TTATGAACCA TACTCTGAGG AGAATCTCAC
AGAAAAGCAG AGACTTCTTT TGGGTTTAAC ATCAGCAGAT GGGAAGTCTG ATAGTTCCAA AACAATTCAT ACTAACAAAT
GCATCTGTCT TCTTTCTCAC TGGGCNTTTT TTTGATGGCA TTCAGGAAGT TTCTGACTTT TNCTGTATCG TTAATTCCAT
CTCTGGGGGCT CATGTCCTTC CAATTGAGGA GGATAATTCC CA

SEO ID NO:371: (Length of Sequence = 441 Nucleotides)

GACAAAGTCA CTCAGGGICT ATTICACCAT ACCCCAAAGT AAAGGCCCAA ACTCCACCGG GGCCAAGINI TICTGGNICA
AAGTCACCAT GICCCCAAGA GAAGTCIAAA GACTCACTAG TICAAAGTIG CCCTGGNICC CICTCTCTCT GIGCAGGAGT
AAAATCIAGC ACACCACCAG GCGAGAGCTA TITTGGIGIC TCATCTCTGC AACTGAAAGG ACAATCTCAA ACTTCACCAG
ACCCACAGATC TGATACTTCA AGTCCAGAAG TGAGACAGAG TCATCAGAA TCACCATCTC TGCAGAGCAA ATCTCAAACA
TCACCTAAGG GAGGICGGIC CAGGICTTCA TCTCCAGICA CITAGCTIGG CATCCAGATC TCCANIAAGG NCAAGATAGA
GGGGAGTTCT CAGCGAGICC TATGTTGAAA TCTTGGAATT T

SEO ID NO:372: (Length of Sequence = 362 Nucleotides)

GAGGIATIGI IGITACIGGG AGGITGAAGG GAACACAAAT TCAGITATAA GICCITITIG AATACIAAGA GGGGAATAAT
TAGGGAAGCT AAGAGGGGAA TAATTAGGAG AAGAAAAAAA AACITCAAAC AATTITCCCT GIAACATGAT TITACITGCA
TITATAAAACT GATTITITIT TCTAAGCACT CCTITGATAA TGATTAAGIG TGGGGITACA TTATTINAGG GICGICTAAT
ATTIAAGGIG ACITAAAAAC CICACACACG TTAATCCCGA ACIGIGAAAA TTICICATCT TATCATCCCT CIGITACTAT
CAATTITCCT CACGGIACAG ATTCTTTIAT AATTACTICA TT

SEO ID NO:373: (Length of Sequence = 306 Nucleotides)

ATTCTTTGTG CGTGTGTGTG TGTGTGTGTG TGTGTTTTGC TGTGGAGTTG AGTTTCTTTG TAAATTCTGG
ATATTAGTTT CTTGTTAGAT GAATAGTTTG TGAATATGTT CTCCCATTCA ACAGGTTGCC TCTTCATTCT GTTGATTGTT
TCCNTTGATC TGCAAAAACT TTTNACTTTA ATATAGTTCT ATTTGTTTAA TTCTGTTTTT CTTACCCATG CTTCTGALAAT

CTTAGCCATA AAATGTTTGC CTAGAACAAT GCCCTGGAGT GTTTCCCCTG AGTTTTCTTC TGGTAG

SEO ID NO:374: (Length of Sequence = 278 Nucleotides)

GGGTTTIGGT TGAGGTTTCT ACCTCATTAT CCAAGATATT INCTITICCAG CCAGCAGAAA GAAAAAGGAG AAGAGCTGCC ACCCTTTGTA TCCAGGATGA TCTCTTNITG AAATCCTTGA TTTAATTATA TCTGCATGAC CCTTTNCCCA ACTAAGGTTA TATCCACAGT TACCGGGGGT TAGCACTGGG ACATCCCTTA TTTTANGAAC ATGICTCAGA AAGTTGCACA AAAAACTTCT ACTACATCCC ATTGGCCAAT ACTTCTTACA TGATGACA

SEQ ID NO:375: (Length of Sequence = 321 Nucleotides)

GGTGACAGTA TITTTIGIGG TITCIGIAGC TCCAGCCCCT CAGAAGGGAC GCCTACAGTT GGCAGCTATG GCTGTACCCC
TCAGTCATTG CCCAAGTTCC AGCATCCTTC CCATGAACTG CTCAAGGAAA ATGGCTTCAC ACAACACGTC TACCATAAGT
ATCGTAGGCG CTGCCTTAAT GGTAAGAAGT GTGGGGGCCA GGAGATGAGC CTCTGGGCCC GTTATTTAGA CCCAGAGTAT
AAGAGTTGGG GGATACGGGG ATAGGTGACT CTTTTCTCTG ACTTCAGAGC AAAAAAAAGA CATGACATTA TAGCAAGAAA
G

SEO ID NO:376: (Length of Sequence = 337 Nucleotides)

GGAAAATTIA CAGCATGACT ACATATGTTA GGAAAAAAAT ATCTAAAATC AATTAACTAA GCTTCCATCT TAGGAAACTA
AAAAAAGAAG AGCAAATTAA ATCCAAAGTA AGAAGAAGAA AATAAATAAT AAAAATTAGA GCAGAGAGAA ATGAAATTAT
GAACAGGAAA TCAATTTTAA AAATAAATGA AACCAAAAGC TGGTTCTTTG AATCAATTAA TAAAATTGAT AAGCCTCTAG
CCAGACTAAG AAAAAAGAGG TAGGGCACAA ATTACTAATA TCATAAGTCA AAGAGGGGAC ACCCCTACAG ATCCCATGGA
TATTAAAAGG ATAATAA

SEQ ID NO:377: (Length of Sequence = 455 Nucleotides)

GTTACAATIG AGAAAACATA TITAATAAAT CATTGICAAT TITINATAATG TITCAAGCCC ATTCITTGIT GATAGCCTCC
ACATTTATAT GGTTAAGICA TIGITGCIGT GITTCTTACC TATGACATTA TITINATATC CCTTCATTIG TGGATCTTAA
GATGITGCAG AAGGITCATT CCTGTACCCC AATACAGATT CACTTCCTTT AGCTGCCTTT NCTAGCACCA ATATGCTTTA
AAAAAAAATG CGCAAACAAC AAGCAGTGAC AGCGGCCAAT TCCTCGAATG TCCAGATTAA TAACTGTAGC ATGCTAAAGA
AAGGIGTGIG TAAATAGCTG GAGATGGTAT ATGGTCCAGA GTCCAGCATA AAATTATTTC CTTTCTGAGG CATTCCCTCC
ATTCCCCTAA CCCGGATACA TGCATTAGGA ATGTAGCAAA ACCCTTCGGG GAACC

SEO ID NO:378: (Length of Sequence = 349 Nucleotides)

GATGGTCACG GGTGTTTATT ACTGGACATG CTCTATGCTT ACTTGCTTGA AAACGCTCCA TTAGAAAAATA AACTCTGAAA
ACTATATGCC CAATGCTAAT AGTGGGTATT TATTGGTAAC ACTCTTTATC AGGTGCTATG ATTGTTGATG GCTTTATTTIN
CINCITCATA TTINCIATAA TINCIACAAT GAACATGTAT GTATAATCAG ACAAAAAAGC CAAGAAATAT CCATAAGTTT
TNCTGGTCAT TCATTCATCC CATAAATACT TGCTGAGCAC CTGCTGTAAG CCAGGCTCCG AGCCGGCTGC TGGGTGGAGT
GCCGCACCCC AGGGAACGGT CAGCCCTCG

SEO ID NO:379: (Length of Sequence = 421 Nucleotides)

ATTITIGAATC ATAITITACT TATAGGITIG CIGIATATAC IGAITAAACT ICIGAACCTA AAGAITCICT ATAATTAAAC TAGCACAAAAT ATAATCIGIC CCTTACCCAC ATIGIAAGAA IGICIGGIGG GGGAAATCCA ATAITGACCT ICACATICCA CAIGGAAAAAT CTITIGICCCC AGAGIGCAAT TAGGGIGATT AAAAATAAGC AGCITITIGIG AGICICAAGT ITIGITCCCCA AALAACAAGCAGC ATCAGCAACT GGAAATTIGI CAGACATGCA AALIAGULAG ICCCACCTGA CACATCIAGC CAGAICITAIG

GATCAAAAAT TITGGGGGIG ACCCIGGGCA ATATGGGCIT TAATAAGNCC CIAGGATGGG TICTGATGCA TGCTCCAAAT TITGNGGATCA TIGNINCINT G

SEO ID NO:380: (Length of Sequence = 311 Nucleotides)

ATTTINAGAT GGAGTCTCAC TCTGTCGCCC AGGCTGGAGT GCAGTGCCAT GATCTCGGCT CACTGCAACC TCCACCTCCC
AGGTTCAAGC AATCCTTCTG CCTCAGCTTC CCCAGTAGCT GGGATTACAG GCACCTGGCT AATTTITTTT TTTTTTTTTT
TTTTGAGATG AAGTCTTGCT CAGTCGCCAG GWTGGAGTGC AGTGGTGTGA TCCTGACTCA CTGCAGCCTC TGCCTTCCGT
GTTCAAGCGA TCCTCCTGCC TCAGCCTCCT GAGTAGCTGG GACTACAGGC ATGCACTACC ACACTTAGCT A

SEQ ID NO:381: (Length of Sequence = 442 Nucleotides)

AATCIGICAA CATATATITI NATITATCIT AAATACCTAA GAGIGAAATI NIIGGITCAT AIGIGGGIAT ATATICAACT
TIGIAAGAAT CIACCAAAAT GAITITCCAA GIATATGIAT AATGITATGG TCATCAGANC TACATGATAG TIAGAGITGG
TIAACATACT CACTGCAATG GAITGACTITI CCTGTGATTC AGCIATCCCA CICTTAGGCG TATACCCAAG AGAAACTCAT
AATGICCTIG TGIGCAGCTI GIATGCTAAT GAITITAGTA GIATTITTIG TAATAGCCAN AAGGIGGAAA CANTGAAAAC
TITCACGGAA AIGATTAATG AATTAACAAA ATATTATATA TCTATATATG ATCCATTAAT CAATGAAANG GANTGAAGTG
GIATACAAGA AACACCACAG GITAACCNIT GAAAGTATAT TA

SEO ID NO:382: (Length of Sequence = 337 Nucleotides)

AACAGACTIT GGAGCCANTC CCATGTGAGT TIGAGTCTCA GAGTGACTCT GGGCAAGINA CTTAGGCTTT CTGAGACTCA
CTTTCCTCCT TTATAAATCA GGAAGAATAA TCCATTGCTC ATTGAGTGTT TAATNAGACA TAAATGAGAT AGTGTATCTA
AAATGTGATT TGTTAAGTCT AATACGNAAT AGATGCCTAT TTGAGTGTTT CTNATACTCA GGATGGTTCT TGGGATATAT
TTNCCCATGG AACAAAAAGC AGACTACTCA TGACCACTCG GATTTTATGT TCAGCCACAT TAGGGCTCTT ATGGCCTGAC
CTGAAGACCT ACCATTT

SEO ID NO:383: (Length of Sequence = 421 Nucleotides)

GIGAAACIGA AGAAGACCAC GACAAACGAT CGCTCAGCCC CTCGCTTTC TTAGGTTCAC AAGAAATGCG CCGGIGGGGA
ATGAACINIT TCATTAATAA AACCIAATIT GICITGATCC ATTCCACTCT ATAATAAAAC AAAAGATTTT NTAGGCAACT
CGGAATATAG CICITITGAA AGTACTCGAC ACCITTAGAT AAGAATTAAA ACCAACCTAT GIAACTGACA TAATCTTGAT
CINTTAATTT GIAAATATTG ACANITINCT TTCTGCACAT TTTAATCTTA GITTCCCTTT TGATTTINCT GAAGGIGCCA
AATTCCATTT AACINCTITA CAAGICTTTG TAAAATTTTA AATGCATAAA GGGGGGTTGG GGGCAGGGGG ACCNCGGANG
TAGTTTAATT TTCGGAAAGG G

SEQ ID NO:384: (Length of Sequence = 420 Nucleotides)

T

GEACTCCGIT CCCAAGAATA AGITTIGCIT GGGCGGAAAG TATGIGGITC ATCCGAAAAA AAAGAAATCA ATGATTIGIG GCAGITCITC ATGIGCITIT GGGCATIINC ATATCITCCT TGGAGAAATA TCAATTAAGA TCCATIGCCG TATATACATA TATIAAAATT ATGGGICATG TATIATGGCT CATACCIGIA ATCCCAATGC TITTIGGATGT TGAGGCGGGA GCVICACCIG AGGITAGGAG TTCGAGACCA GCCIGACCAA CGIGGIGAAC CCIGICTCIA CIAAAAATAC AAAAGITAGC CAGGCATGGI GGCATGCACC TGIAGICCCA GCTACCCAGG AGGCIGAGAC AGGAGGAATT GCTIGAACCC ANGAGGCAGA GNITICCAGT GAGCINAGGA TIGIGCCACT

SEO ID NO:385: (Length of Sequence = 404 Nucleotides)

GIGACAAATG TIAAGAAATT GIGIGICAAG CAAAATACIT TAGAGGCCAA TGGGCCACAT GITTITAATA TCAAGAGATT ACACACAAAA TIINITITCI AGCITCITIT GAAAAATCAG AATIGGGAAG ATGIATICAT GAGIGACIGC TGCCCCCTIT GGITGGGACT CGITCCITCA GGITCATTAC ATGGCCATCA ATAACCATTI CCITGGICCC TGCITTIGIC TIGICIGGAC TCTAAGCATT TGAATTITTA GIATTATAAG AAAACITAAT ACITINCIAT CAGICACCAC ATACATGIGI TICIATCIGI ACTACGACT ATTAAAAAGCN TITIATCAAT AGCCNCCATT TTGGAGGGGG GGATTICAAC TGGIGCCING ACTAGCAAGG AATI

SEO ID NO:386: (Length of Sequence = 267 Nucleotides)

GICTIGIGGA CATTIACGIG GIATCITTAG AGCAAACACA GAGIGGITGC ATAAGCIGCA GIGITTIAGI ATCOGIGOGGA CIGITGGGAG GCGIAGAGGA GINACAGICG CAAACIGAIG GCCCAGCICT GACCCTCCAG GCAAGIGGAC TCCGAGGAGT ACCAGCAGAT CITCCCACAT GCGICGGGGA GGGCICTGGG GAGAGICAGI GGGCAGGAGA GGGTCAGCIG TGCAGGCICC AGGGCCCAGC CCCGIGCTIT CCCCTCT

<u>SEO ID NO:387:</u> (Length of Sequence = 384 Nucleotides)

ATTITAAATG ACATTITATT TAGGCCAGG GACCAGGTAA CATTATTTTT AGGAGGAGAG CAAAAGGTGT TATATTACTG
CTTCTAATTA CCTAGAAGGA AAGCATTTGC TACACTGCCA TTATGATTGG CTGCAGCAGT TCAACCTGGC TCTCGGAATC
TGCCATTAGC TTGACAGCAT ACAGAGCACC ATATCAGGGT TACTATGGGA AGACTCTATT GTGGCATCAG AAACACAAAA
AACACTGGAT ACAGTTAGTT TCTGTTGACA GTTTCAGAAG AAAATCCCAC AGATTGGACA GGCTGCCTGC TGAAAGGGTT
GTCACTACAC ACAGCATGCC CTGAACCCTG GGAATGAAGT TACCCCTATC TGTGGTGATC AGGA

SEQ ID NO:388: (Length of Sequence = 345 Nucleotides)

CTAAGATCAA ATGCAGGCAA AAGTGGTGAA TITTACCACC TGTTTGTAAG TCTGGGTTTA TAACTTTACC GTAAATCACC TAGAACCACA GCTAGCCGAA TCGGGGTGC TGGTATGGCA ATATCCCGAG AGCTAACCTG GGGCTGGGGC AATGTTCTGT GGCTGCTGCA CTTGCCTCTA ACAGGCCAGT TTAAAACGTC CAGCTCTCAG GGCCACATTC TCCAGGACAC AGCAGGGAGC TCACAGTAGC TCAAGACCCG GCCCAGCCTC CATCCCCAGC CTTGGAGCTG TCAGTGCTCC CAAAGGCTGA AAGAATTCGG TCTTGGCTGA GTGGACAGCC CCCTT

SEO ID NO:389: (Length of Sequence = 156 Nucleotides)

TAACCTGCCC CAGCAGTGCA TGCAGGAAGA CITCCTGGTG CATGAGGTGA CCAATCTGCC GGTGACAGAA GNACTGATTG AGCGGGAGAA TGCAGCCCAG CTCAAGAAGT GGCGGGAAAC GCGGGGGGGCC CTGCAGTATC GGCCCTCACG GCGACT

SEO ID NO:390: (Length of Sequence = 364 Nucleotides)

GASTICICCCT CIGICACCCA GGCIGGAGIG CAATGGCAIG ATCICGGCIC ACTGCAACCT CCGCCTCCCG GGITCAAGIG
ATTCICCTGC CICAGCCTCC CGAGIAGCIG AGATTACAGG CACGIGCCAC CACGCCTGGC TAATTTIGIA TITICAGIAG
AGATGAGGIT TIGCCATGIT GGCCAGGCIG GGCTCAAACT CITGACCTCA GATGACCCGC CIGCCTCAGC CICCCAAAGT
TCTGGGATTA CAGGCATGAG CCACTGCACC CAGCCCAACA CIGGGATTCT TTTATCCGCT GGCTGGCTCT TCCGCAGTTG
AATTGTGTGA CTTCTTCCCC TATCTGAGGC CCAGTTTTC TTCA

SEO ID NO:391: (Length of Sequence = 325 Nucleotides)

GAGTGTCCAG ATGATGCCAG TGATGCCCCA TCTGGAGCGG CTGCTGTAAG GACACTGGCT GCAGCAGGGG AGGCACAGCC
AGGCCTGCGC ACTAGGCAGA GCTGGTGTGG GAGCCAGGAG CAGA/GALAG CCCCGCCTTC TACCAAGTTG GCALTGCAGA
AGGCCGCCACT CCCGGGTGCT GATGCCGAGT TCAGGTCCAC ACCTGGCTAT CCCTGGCCTA TCAGGGGCCC AGGAAGCCC.

CCACCCCTGC AGENTICAAA GGGCCTGCTT CCCACTCCTT GGCCTTTCCC TCCTCCTGGG AACCATTCTG GGGCAGAGCA

SEO ID NO:392: (Length of Sequence = 371 Mucleotides)

ACATCCACAC AAGTACAAGA ATACAGAAGC TICTCTAGIC AGGATGCACT AAGCACCTAA TGAGTAAACA AACTICAGCA
TATCCTCATT GITCTCATGG TATTAATTIG AAGATACTTA CCTTCGAACT AAATCTGGIT TTAGAAGAGC TGCTTGTTGT
TCAGCTCCAA CTGGTTGGGA TACAGGCTGT AAACAGTACA GACATAAAAC TTGCTATGAT AACAGTAAAA TTCAAGCTAA
ATATACAATT TGTTACTATT CAGAAAACAC GATAGTTTTG GTTACCTTGC AAACCTGGTA GGAATATCTA TGTTATTGAA
TGTCTGTATC AATCCTATTA TTAACATTAT TACCAAAGGT AAATAAAATT T

SEO ID NO:393: (Length of Sequence = 404 Nucleotides)

CCTITIAGIA GCITCICIGA GGIGAAACCA CTTCTITITG ACCATCIAGC GCANICINIC TITIACATCAA CCATITATITI CAAGIGIAGI GIGCITCAGA GICIGAAAGA GCIATIGCAG AATIGGCIGI TGIGGCITTC TAIGGACATI CACATGAAAC CTGITIACAAA CAGICCICTA GAGACAACIT TGGGGGATC CATGAACICT GIGICTAAAC TGATCCACIA TGIAGGGIGG CTATCCACIA CTGCAATGCG CTTGGAGAGC AACAATACIT TCTTGCTGCA CTTTATTTTG GATTTCTATG AGAAGGIGTG TGACATATAT ATAAATNATA ACCITCCATT AGIGGGIATT GITTCCTCCT GGGGATCCTT CTATTCTGCA CTCCTCAGCC TGGG

SEO ID NO:394: (Length of Sequence = 416 Nucleotides)

GCCACACACT GGAGAGGGAG AGCTAGAGAG TGAGACAGCA GGGGAGCTGA GGGTGAATGG CTGCTGTTAG AAGCCCTGGA
GACAGCCTGA GGTCAGAGCC CAGCCCCACT CCTGGCTGT TGATCTTGAG CAGGGCTGTT AACTTCACTA GGACTTGGTT
TGGGTTTCTC ATAGAGAATA GGTACAGTGT GAATTAAATA TATATAGCTT GAATAAAGTG CCCAGCTTGT GGGTAGCTGC
TGCCATCATC ATCACCATCA CCATCATCAC CATCACCATC ATCATCATCA TCATCATCAT CATCATCATC ATCATCATCA
TCTCAGGCAC AGGGGCTTTA AGGACAACAT GCCCAGTTTA AGGANGAACA CAACTCTCTT CATTTATAGC GNCCCTCCAT
CAGTGAGTAG ACGCTT

SEO ID NO:395: (Length of Sequence = 315 Nucleotides)

AGAGATCAAA TGTCTTAAAC ATTATGGAAT AGGAGTGTAT GACTGACTAA CATCCAGTAA TCATTAGGGA AAACAAACAT GAGTGAGGNC AACTGAAATA ATTATGATAC AATTAAGGGT GGTAGGTTAC ATTTGTATAG TTCTTTAAAA TATGCATTAT TCCACATGAT CAGAAATATA AAANGANCTA GACAGATACT GGTAGAGAGA CAATTAATTT AAATTTGTAA CATATTGCTT GGNGCAAGCA TTCAAGTTGA GTGCTTAATG TGTATCGGTG ACTGCACTGT GCAAATAAAT TTGGGGTTAG TAAGA

SEO ID NO:396: (Length of Sequence = 409 Nucleotides)

CTCCAGITCT CACGITAGGG TECTITCTIC CCCGGCAGAG TITITCGAGC TCATGAAGGI GGACIGCCIG GAAAGIACTC
TAGAAAAGIC ACTCCAAGCA AAGITTCCIT CAAATCTCAA GGICICCAAT CTCTTAGACT TCACGCGGGG CTCACGAGGC
CGGAAGAACT CCCGCACAAT GCTGCTCCCA CTCCTGCGGA GGITCCCAGA GCAGGTCCGA GTCTCCCTCT TTCACACGCC
GCACCTCCGT GGGCTGCTTC GGCTCCTCAT CCCTGAGCGC TTCAACGAGA CCATCGGCCT CCAGCACATT AAGGTGTACC
TCTTCGACAA CAGCGINATC TTGAGCGGTG CAAACCTGAG TGACTCCTAC TTINACCAAC CGTCAGACCG NTACGTGTTC
CTGCAAGGA

SEO ID NO:397: (Length of Sequence = 414 Nucleotides)

ACAAGCTGTG TGACCATAGG CAAGTTGAC CTITCTGAGC TGCCATTTTC TCATGGTAAA AGAGAGATAC TAGAGGAACC
TGCCTCACAG GATTGTCATG GAGAATAGAG GAGATGATAC AAGTGAAGCA CTAGGCAGCA CCATACTTGG AACTAAGGGA
AAGCCCGCAG TCAATGTTCA GTATTGTTAC ACTTGCCAGA TTGTGAAAGA GCGCAGGCAA CCCTTGAGTT GAGCTCAACG
CTGGAGCCCAA GATCAATGAC AGAAGGATTT TGTTTTGAAA CAGCAACTAA TGACCAGAGA GAGGAAATGG GTCATGAAGC
TCCATGGTGC CTTTCATGAA AATGAAATGT AAGGGCGTGA TTCAGGAAAA AGGGACCACG ATCAATACCA GCAGACTCTT
CCCTTATGCAC TGGG

SEO ID NO:398: (Length of Sequence = 400 Nucleotides)

CATCAAGCTG GGAATGCCCT AAAGTGGGGG CGTGAGGAAG AGAAGGGGTG ATACCTAGAG GCTGGGGTAT CTCTGTCCCA
AGGAGACAAA CTATAACAAG ACCCAGCAAC TGAAGGGTTA ACACCTAGCA CAGACGTATA CCTCCAGGAT CCTAGCTGCA
TTTCIAATTC TGCTTCATCT ATGCTTGAGC ACTACTTGTT GTTAAATATA CTTAATATCA CTCTTAGCTA ATTTCCTCTA
TGTAGATTTT TATTTATTTC TGAGGGCAAC CCAACTTCCA GGCTCTTGGA AGGAAATAGA CTGCAGCCCC TAAGTGTGAT
CAATACTTAA TTATAACAAT AATCACTAAT AATAACTTGT GCTGCTTCAT TGTAACTAAA ATGTACACTT TTACATTTTT

SEQ ID NO:399: (Length of Sequence = 324 Nucleotides)

AAATATTTAC AATTITACAC CTTCAGGAAG GCTCCAAAAT ATAAACACTG TACCTCTCCC TAGAGAAAAA AAAATTATTC
TTCTCTTCAA AAACAGGAAT ACATTCATTT TTTCTCACTG TGTGAATCAA GTAATTATAC AAATAAACAT CTGAAACATT
TTCCTTTTTA ATATATTAT ATAATATATA TTTMTAACAG CTTTACAAAT AAAGGCAACG GTCCTTTTCT AATTITCATG
CCTCTCAACA GAAGGGTACA TGATGCTCCC TGAATTCCAG GGNTATTTTT TNCTCTCTAT GGTACTTTGT ATTITCACTTT
ACTT

SEO ID NO:400: (Length of Sequence = 388 Nucleotides)

ATTAAATCIG AGITTIGITT GAGCATCTIT CAACATGTAC CATATITATG ACAATTCCT TCCATAGGAT CTATCTGINC TGCAACAAGT ATTGATCTTA CAGTAAAATT TTTCACAAAT TCATTAGATT CTATGTCTCT TTTTCTGGTA GGAATTTTTG TGCAGGTAGC TATCTCTTGC CCTAGATTAT TCTCCTTGTT TAGCTGCTGA TTCTTAAACT GGCCTCTAGA TTTCCAGATT TCTTCCGGTA CAGACTTTCT CTTTGCAAGT NCTTCCATCT CTAATCTTTG AGATTAATCT TCTTTTGAAA TGTCCTGCTG CTCTACTCTT GTATGTCTTG GNCCCACGTT CAAGCTTCCC ATCTAGCAAA ACCAGGGTTT CTAATATT

SEQ ID NO:401: (Length of Sequence = 339 Nucleotides)

GITTATTECT CAAAAACAAG AATTCAGAAG CAAAGGTEGA GAGACTGTGG GITGGGGAGA TGGCAGGAAG GGGGCAAGGC
CITGTCCCAG CTCTCCCCTT TGTCCTTCTT CTGACCCTCC TGGCCGGAGT CAGGCCTAGG GCCAGGGCAT CTGGGAGGGG
GGCACCTTCG TGGCCAAGGG AACAGTAGAG CTATCGGGGG CAGTCCTTGA GGGGTGCCCT GGGCAGGAGG GGCTGCAAGA
TTTNCAGGGA GGCAGAGTTC CCCTCCCAGA ATCCAAAAAGC CGGTAGGGCG GGGGGCAAGG CCCCTCGTTT GGCAACTNAG
AAGAGGCGGC TTTTGGGCG

SEO ID NO:402: (Length of Sequence = 400 Nucleotides)

TGTCCAGTGT ATGAGGACGT CCCAGCGAGA AATGAAAGGT TCTATGTTTA TGAAAATAAA AAGGAAGCAT TGCAAGCTGT CAAGATGATC AAAGGGTCCC GATTTAAAGC TTTTTCTACC AGAGAAGACG CTGAGAAATT TGCTAGAGGA ATTTGTGATT ATTTCCCTTC TCCAAGCAAA ACGTCCTTAC CACTGTCTCC TATGAAAACA GCTCCACTCT TTAGCAATGA CAGGTTGAAA GATGGTTTGT GCTTGTCGGA ATCAGAAACA GTCAACAAAG AGCGAGGACCT CACCGCCAAG CTTTCCGGAA AGCTGTNGAG GAAAGGGAGG AGGAGGACAN CTTTTCTGAC CTTATCTGGG AGCAACCCCC

SEO ID NO:403: (Length of Sequence = 416 Nucleotides)

AGITGACTGC TCTGATATGG AGAGACCTGT TAGICTTGTA TATAGTGCCC AGCCGGAAAA AGCATCTCTT GAAGGTTAGG
GCATTTTGTG AGGAGACCT TAGGGCTATA TCAGTCTGGA GGTATGATCT CTGATAAAGA TCATAATTCT CATCTCAGTA
ATCTTCTTTA GAACAAAACA TTCTTCATTG TAAGCTTCTC ATTAACTGAA GGCCACCTGA TCTGAGATTT TGGCTCTTAG
AATACTCTTT NCTGTGTCTC AATCCTCATA TGGCTTACCT CTGAAATATA GAATATATTT CCTTGTGTAG CCTGGTAGAG
TTGGGTTTTG TTTTGTTTTT CAAACAGTAA CTTTTATTTG ATTGTAAAAC TTCCAGATTT CTGAGATGCC GCCTTACCAG
TCTTAAGGTT GATTTT

SEO ID NO:404: (Length of Sequence = 368 Nucleotides)

CCICINACIC ATTGIGATGA GIAGGGGGA GGGCITCACI GCCICANITI CCCCAACITI GGACCITAAA TCCICICCIG
ATGCCICCA GCCCAGCCAG GAAGGAGAC TAAGACCAAG AGGGATTAA CAGATGCAGG ACACACAGCC TIGICCICAG
ACCCCCCAAG TCTGAGAGAA GCAAAACACI CACCITGAGA GCCCICGGAC TIGGAGGTGA GGIGCAGAAC CCAGGCIGGG
TGIGIGCIGA GGGGIGGIGG GGGGGGGGG TGCTGGGGGGG AATACITITC TTAAGCTAAG GCIGGGGCTT
AGGGGAGGGC CAGAGGAAGG GIAAATAGIT TGCCIGGGGG GGIGCIGG

SEO ID NO:405: (Length of Sequence = 395 Nucleotides)

GACAGGICCT CACTCTTACC ACAAAGCTCA AGTCAGCTTG GCCTCTCAAG TGGAGAGGATA ATCGITCTAT AGCAAGAAGT
ACAAAGATTC TCTGCAGACA AAACCAGCTA GCCAAGGTTC CACAACATGT GTACACGTAT AAGTCTGNTG GATCAGAAGA
AATATGTACC CGGGAATCAG ATGTAGCCAG CCCACATACT AACAAACATC AAAGCAAGCC TAGTCAGATT GAGTCCCATT
TGAACAATCT TTATAAAGGT TTCTTCATGT TATTTACAAT TCAAAGTAAA TTTACTTTAT AAGCAGCTAG GGGAATTCTT
TATTTAGTAA TGTCCTAACA TAAAAGTTTC ACATAACTGG CTTCTGTCCA AACCATGGAT ACTTGAGCTT TGTGG

SEQ ID NO:406: (Length of Sequence = 358 Nucleotides)

GATACCITAA TCTAAATTT ATCITAATTT TEATITITAT TICATEGICT AAATTITIAT CIAAAATTTI INCIAGCICT TIATIACACC AAGACAGCTT CACATITITA TITATATATI GIACATCICA TGIAAGGNAT TACCGIATAT AAGCTAGIGT CATAACITAA GIAGCCACAT TCATTCAGIA TGITTTATGT TITCICTCTG ACTGGATCTC TGATACATTC TITCCTGITC TAGCCIGCTT TATGCAAAAG GGCATTATAT GITTGICAAT CAACCAGGCT TCTGTGACTG TTTAGAAAGGA ATTATGTAAA TATATAATCC NGIGGCCTGT TTCACTTTGG CCATGTTT

SEO ID NO:407: (Length of Sequence = 294 Nucleotides)

CIGIGIATAT TTAGIATCIT TNATTAAGAA GACTGGITGA TATTIGCCIT CAGCIAATIT ATAGAAAGGA TGATCATCAA TGICICIAGI TITCITCIAA GIGGCITGIC TGIGCAGGIA CATATAAAAA TNCAACIATA CAAATAGCIG GACAGITGAG TCICAACIAT GAAAATCITT TCIGGGATCA AGATCIAAGA AGITGGIGIG TGIATGAGIG CAACCCATCA TTCIATCCCC TAAAAAATCIG GGGITTCICA GCCCAAACAT TCNCACIAGI AAAGICAAGI TTCA

SEQ ID NO:408: (Length of Sequence = 367 Nucleotides)

10

SEO ID NO:409: (Length of Sequence = 233 Mucleotides)

AAGAGACAGG GNCTCATTCT GTCAACCIGG CIGGACTGCA ATGGTGTGAT CACAACTCAC TGCAGCCTTG AACTCCIGGA CTCAAGCANT CCINCCACCC CAGCCTCCTG AGCAGTTAGG ACTACAGATG GGTGCCACCA TGCCCAGCTA ATTTCTAAAT TTTTTTTAGA GACAGGGTCT TGCTATATTA CCCAGGCTGG TCTCAAACTC CCTGGGCTCA AGTGATCCTC CTT

SEQ ID NO:410: (Length of Sequence = 295 Nucleotides)

GACAGGGGGT GGGGAATTCT ACTOCATGGT ATCTTCAGAG CTAGGATAAT GCTCCTTATG CAATCCCACT GCATATGACC ATGGCAGTAG AACAAGTTCA ATTACTACAC TGGATGCGTT AAGTGTGCTT TCCTAGCAGA AAGCACCAGG GTGGAGTCAA CAGTTCACAT GCTAATACTT GGAAGTATTT CTAGAAGGGG GTGCTCAATA GAGGGCAGAC ATGATGCAAG NNCTTCATAC TAGAAAGGTG TCCTGTGTGT GCATGCACAG CTGGATGGGG GCACACAGGA GCAAG

SEO ID NO:411: (Length of Sequence = 304 Nucleotides)

AATAAAAGA CCATTAACTT AAAGTGGTGT TAAATGCTTT GTAAAGCTGA GATCTAAATG GGGACAAGGC AGGTGGAGGG
GAGGCCAGTG TACATGTAAA TGCCCACAGC CCAGCATTGG GTTTCCCTCC CAAGGNCCCA GCACCAACCT CTGAGCCCAA
GACCTTGCCT GAAAACAAGC AGATACCGAT TGNTTCATCC TATTTATGGA CATGTAGGTC TAGTTGCATT TTCACTNGGG
GGAGGGGGGA AGGTGAATTA TGGTAACTTT TAATGATCTA TTCAGGCAGT AGAGCTCTTA AGGG

SEO ID NO:412: (Length of Sequence = 250 Nucleotides)

CAGGIGOGCA CIATCACGOC CEGATAATIT TITTIGITIT TAGTAGAGAC GEGGITICAA CATECTECTC AGGCIGGICT CAACTACCGA CCICCIGATC COTCCACCGC GGCCICCCAA AGTGCIGGGA TCACAGGCGI GAGCACCNCT CCIGGNCACA GGINGAGACC CITTCIATAT AAGAAAGAGA AAAATGICTC TNANICACAA GAGAATGCTA ACAACGGGGG AAAGCACAGA CACAAACCTG

SEO ID NO:413: (Length of Sequence = 337 Nucleotides)

GTACTIGGGAC AAGGGAAGGC AATCACAAAC AACTIGCCCTC AGGAAGAACT CAGTCCCTGA CTGTAGTGTC TCTTCGGGGG
AACCAATGCC ACCCNCCTCC ATCCCCCAGA CGGGCGAGGG GCTGCACCCT TAAAGCAGGC CATTGGGCCT TCCGGGCTCC
AGGGCCAGCC CACCCCCGNTC CCGCTGGTGG ATCTTCTGGT GCTGCAGGAG GTGCTGCTTC TGGACAAAGC TCTTNTCACA
CTCAGTGCAG CTGTAGGGCC GNTCACCCGT NTGGATGCGC TGGTNCCGNA CCAAGTCAGA TGGGTGACTG AAGCTCTTGC
CACAAGTAAC CACAGAT

SEO ID NO:414: (Length of Sequence = 304 Nucleotides)

GGITTAAGAA CIGCGITTIG GNGCCCAATC TITGGIGAAA AATATITTIG GGICATCITT GAAAAAAATC CITTTCAAGG
CAGACAGCAT TITAATGCIT TGICTGITTI TCCCIGITTIG TCAGCTCTGN CACCAGCCTG AAAGATTTAA AAAINCAAAT
TAATGGAGGN TTATITGTCC TNIACTCAGG TCACATTTCT GGGITTTAAT GAAGNGACAG ATGCTGCTCA TATACAGGAT
TTAGCTGCAG TTTCTTTGGA ACTTCCAGAT ATTCTGAATT CACTCCACTT CTGCAGTCTA AATG

SEQ ID NO:415: (Length of Sequence = 315 Nucleotides)

CGTTGTGGAG TGGGTGTCTT TGGATAGAAG GAGTGAGGAA CTGGGGGAGG AAGGCCTGGG GGATCCCCTG GCGGGGCTAC TTCCTGGGCC CGGNATGGAC ACCTGCNAGC TGCTGCCNTT GTTGGGGTCC TGGCAGGGGT GTGGTGTGGC CCTCACCACT CTGNTCACCT GCTCCTTCCT NACAGTGCCT GGAGAAGTTC CCTGTNATCC AGCACTTTCA AAGTTCGCNA GCCTNCTGCC CATCCATCCT GTCACGTCGG GCTAGGAGGG GNCAAGCCGA AGAGCCACCC ANGNCACANT TCCTGTGCCT GCCTT

SEO ID NO:416: (Length of Sequence = 343 Nucleotides)

GTATTICAAG TGITTIATIT GCTITCIGIG GIGICAAATI TGGGGTCTCC TAGAGCCCAG CCCCAGGCAG AATCCGGCAT ATCCTICTCC GCCTGGGGG CCCGGGACAC AGGAGITTCA GAAAAGGCAC TGGCAAAAGI NCTAGGGCGG GGGTCAGGGA GAAGCCACAC TGAGCCTGGA GGGACCGGGC CCTCCTTCGG CGGCAGAAAA CACAGTCACC TITNGCAGGG AAGGGTTTITI NCCTAGAAAG AAATTTAAGA CAAGATAAAA ACCTGAGATG TTAGAGGAGC CCCCAGAACC AAGCCGGTGC TNCCCTGGGC AANCAGAGAG TGAACTCGGC TTT

SEQ ID NO:417: (Length of Sequence = 202 Nucleotides)

TATTICICIG IGAAAAGGGG GAAAATAAAA GGAATAAAAT AAAAACGGCA CAGITGACAC ACAAAAAAAA ACCAATGAIG GGGAGGACGG GAGGIGGAGA AGTAAATGGG GGAGGGGNIC CCATTACAGC AGCAGGATCC AGINACCCGG GAIGCICACA TCINNICCCIN ACGIGGGCGG TGIAGCCCCT TCCICCCAAG GT

SEO ID NO:418: (Length of Sequence = 299 Nucleotides)

CACCAGITIGG CIGCAGAGCT GICTICAGGA TCATAGGCCA CIGCCAGAGT CITGGAGAGA GGGAGAGATG GAGAGGAAGG GAGTGAGCTT COGIGGICIG ATTICIGGCT CAACGACGCA GGAACCTCAG GITCAAAAGC AGCIGACAAG AGCCCAGAGA CCGICTICTT GGCGICCGGC AGAGCCITCT GGIGGCCCGA CACCCAGGCA NGGAGGGAAG GCCCTGAAAT CCCGITITIN TGGCAAGATT NGITTCCAAG AGGAGATAAT GGCTCAATTT TGICTICCCCA AGITGATCA

SEO ID NO:419: (Length of Sequence = 223 Nucleotides)

ATTGITGGGA AGGIAACATT TTTCCATGGI TTINATITIN CCCAAAAGIA TTIATGIAIT GATTIATTIG GNICIGACIC AGGCGACGIA CIGIAAGACG ATATTACTIT AATCATCITC ACATCAGIAT TIATGGAATA GCCACAGGIG CCTCATCCIT TAGIAGGAGI TAATTATACA TTINCIGGCC GAGIAAACAT NICCGAATGG TATGIATGIA TIT

SEO ID NO:420: (Length of Sequence = 406 Nucleotides)

TITIAAATATT AAGITAAGIA TATAACTIGC CCIATGCCAT ATTGCTITAA TCAGGGGACT GAGCATCACA TITIAGATITG
ATGAGTITGG GAAAAGITCI CAAACATCCA GACCCATGGA CCITAAGAAT TACIGCAGAA ATCTCCITCA ATATAGICAT
AGGGAGCATT AATGCTITTG TGGIACTAAA CATATITTTG AGCTIAGATA CAAATCCTTC TTGTCCTGAA CTGATAGGGT
AGGAATTGIT TAGGIGCTTC AAATCCAGAT CTTTCAGGGG TTGCCCACCTA AACTCATCTT TATGAGTAAC TCTAGATAAT
AATACACTTT GGIATCTTCC AAAGIGCTTA TCTAGGCATG GAAAAGITCA GTAATTATCA TGAGGACCTG TTTTTAGGTT
AGGICC

SEO ID NO:421: (Length of Sequence = 281 Nucleotides)

ATCCAGATTA CIGACTIGIA CACAATGGAC CATATGINCT GICCAAAATA CACCIACATT ACACTGIGIG GAACANGAAC CIGGGCITTG CAAAAAAGAA TITATGATTA AAATGTAACC CCCCCCAAAA AAAAATGAAG CITAGAATTA AAGGTAGCCT TITACCCAGA TIGITCACCA GNITGIAAAA TICTAATATG GGICATTAAC TIGITCACAAA TAATTCATAT TIGGNCITAT GGITTTAAGGG CICCAGATTG AAAAGGTGCT CIGAACTICT G

SEO ID NO:422: (Length of Sequence = 220 Nucleotides)

THIGHATHT TAAHAGAGAC GOOGHTHIGC CATGINGCCC AGGCTGGITT TGAACTCCTG ACTTCAGGTG ATCTGCCTGC
CTCGGTCTCC CAAAGTGCTG GGATTACAGG CTTTAGCACT GTN_CT31C1 GCCTGGCTGG CTGGCTGGCT GGCTTTCTTT
CTTTCTMTTT TCHNCTCTC TCTCTCTCTC TCTCCCTTCC TTTCTTCCTT CCTTCCTTCC

SEO ID NO:423: (Length of Sequence = 391 Nucleotides)

CTGTCTCTTA TCTGGGCAG CTTTAGACAT ACTAGCTTGG TTGGAAACTG ATATTAAAAG CCTAAAACAT GTAACTTTNC
TTATCAGGTT ACTATCATGG GGAACTAAAG ATTCCTGGTT TTTTGTATGT NCCATAACTA TACTTTAGTA AGCCCTGATA
TACGGTGTTA ATTTTCCTNC AGTGAAGGAA ACATGAAGAT ATATTTATGT GCACACATAC ATATATATGT ATATATAACG
TATATTCAAA CATGCACTCA GAGGAAGTTA GGGAGAGAAG TTTCTAGCTA AACATGATCT TGTGAAATTC TTCCATATGT
GGAAAAGTCG TCAGTTCATC TGACATAGAG CAATACCATA CATATATACA CACAGGGTGC TATGGTATAC A

SEO ID NO:424: (Length of Sequence = 379 Nucleotides)

TOGGGAGCCT GAGGCATGAG AATCGCTTGA GCCCTGGNGG TGGAGGTTGC AGTGAGCTGA GACCCCGTCA CTGAACTCCA
GCCTGGGTGA CAGAGCAAGA CTCTGTATCA AAAAAACAAA CAAACANACA AACAAAAAAG CCTATTATAA AACAATAGGA
AATGCTGAAG TCTAGTGCAC CAAGACATAC TGAATTTCAA ACTAAATAAA TTAAAATTAT CATGTACATT CCACTACATG
TCAAAACAGG AAAANCCATA GTATTATAGT TGATATGAAA TGANGATTAC ATACANCAGT AATACAGAGN AAACATGAAG
CTGCTTATATT TTATTTGGGN ATAAGGNCAN CAGGGGCCCAA TGATTTTCAC TGCAGATGT

SEO ID NO:425: (Length of Sequence = 448 Nucleotides)

TCCACAGGGC GGCCTGGGGT CTGGAGATGG GCGCTGGGCC CACGGGACGC AGATGGGGCC AGGCTCTGCC CGTGGCTGGC CCACGTTCCT GGTCTGCAGT GCTGCCTCT CCCCAGCACC CCTGGGGCAC AGAGGGCAGG GTCACAGCTG GGAAGAGGTG GGGGGTTAGAA ACCAAGGCTG GCAGAAGTNT AGCCGGGCTC CCTGATAAAT GCTGGAGGAC CCCAGGGCAC CTGCACTTAC TGTACCCTCT CTGAGAGCAT TTGTATGATC TCATGTCTCA GCTCTNNGAG GCTGGAGGTC CCAGAAAACC AAGGTATGGG TAAGATTCAG TCTCTGGGTG AGTACCCAGT TNCTGGCTTC TAGATGGCGC CTTTTTCCCT GTGTGTCCTC AAATGATTGG ATGAGGCCCAG GGTGCTCTCT TGGAGTCCTT TCTGTAAGGG CAACTGAT

SEO ID NO:426: (Length of Sequence = 417 Nucleotides)

GCCTGGNTCA TCGCTGTCCT TTCCTCCTTG TCAGAGTCAG TGACACTGAC ATTAAGGTCA TCGAATATCA ACCAGGTCCT
GAGGACCTTG GTGTGTTTCC TCCTCTCCTA GTCTCCAGAC CCCAGCCTGT TCATTCCTGA GCTTCCTCTG GCACCCCTTC
CTTGGGGCCCA AGCCAAGTAA GAAATCAGCA GGCCCAAGGT GGTGCTTGGG AGGCCGGGGC AGTGCCAGGG GCAGTCCTCA
TACCATCCTC CCACTGGCTT CCCTCCTGCC TGCTCTTAGC CGCCACACAT ATCTCAGCTG TCGAATCCGA TTAGGGNTTC
TCNCCAGTGA GCCAGACAAG GAGGCCACTN GGCAGGGGAG AGAGAGACAA GGACGCCAAG CAGGGATTGG CAGAAGGAAG
GTGGAGACAT GGCTCAA

SEO ID NO:427: (Length of Sequence = 317 Nucleotides)

AACCCTGTCT CTACTAAAAA TACAAAAAAT TAGCTGGGGG TGGTGGTGGG CGCCTGTAGT CCCAGCTACT CGGGAGGCTG
AGGCAGGAGA ATGGTGTGAA CCCAGGAGGC GGANTTGCAG TGAGCCGAGA TAGTGCCTCT GCACTCCAGC CTGGGTGACA
GAGCGAGACT CCGTCTCAAA AAAAAGGGCT GATAATGATA AACAGTGAGC ACTCCGGTCC TTTTTCTTAC GTTTTCCTTT
TTTCCTTCCT CTCCACCCCA CAAGTTTTGC TTTTTTAACCA AGGTGTCTCT GCTTGATGGA AATTCACATG CTAGTCT

SEO ID NO:428: (Length of Sequence = 296 Nucleotides)

GTAATTACAG TATTINCACG TAGAGACGGG TITCTCCATG TIGGTCAGGC TGGTCTCGAA CTCCTGACCT CGGGTGATCC
GCCTGCCTCG GCGTCCCAAA GTACTGGGAT "TACAGGTMTG AGCCACCGTG CCCAGCCGGT TTTTTTTTTT TYTTTTGTAT
AGCAATGGAA GAATGGCCTC GTACACACGN TAGAGTGGAA AGTCCCAGGC ACCAAGGNTT CCCACCCTAC AAGCAAGGTC
AGGGCTTTCT CTTCATCCTT CCAGGGAGAG CACTGAGAGA TGATGGGGGG TTGGCA

SEO ID NO:429: (Length of Sequence = 422 Nucleotides)

GAGGGTTGGA GACAGGAGAC AGTGGGGTGG GAAATCCAAA TCTCAACTGC TTTTGTACTG TCTCCTGCTC CCGAGTGCCC
CANAGCCCAT GCAGACCCTC TGCTGTCTAT GATATCCTGT TCAGCCCTCA ACTTCTCTA CCATCCCTGC AACTGGGGTT
CACTGTGAGC CAAACCAGTT TGCTTCTTGT TTTCTAAAAG CAGGCAGCCC TTCAGGACTG TNTCATTCAA GGCATTTCCC
ACCTCTNTTC TCCACTCATA TCCCTTCCCA AACTGCCTTT CCTCATTTCT CCGTCTCCAG GGAGAGGGAC TNCAGGCTAC
CACAGNCAAA AATGGTGGTC TTCAGTCCTA CGTAAGNCAA NCTGTGTGAGG TGTGTAAGGA CTNAGGGTTG CTCACAAGGG
GACACACAGA NGTGGATGCC AG

SEO ID NO:430: (Length of Sequence = 332 Nucleotides)

CECCATCACC ACCCEGGACA GCCCCACCGC CCACETGCAG GGGNTGGGGT CCGGGGGGGG CTNGCGCCTC GGCGTCTCCC
GGNAGINTCC CETCCAGCCG TCGAGCAGGG TECTTGANIN TNTCTGCAGA AAAGACTCTA GGACCCGGC ACCATGTTCC
CGGAGCCCCCC AACCCCGGGG CCTCCATCGC CCGANACGCC TCCCGACTCC AGTGCCATCA GCCACGGCCC AGTGCCCCCC
TGGGCCCTGG NCACCATCGT GCTGGTCTNA GGCCTCCTNA TCTTCAGCTG CTGTTTCTGT CTCTACCGGA AGAGCTGTCG
GAGGCGGACA GG

SEO ID NO:431: (Length of Sequence = 413 Nucleotides)

TGTCATTATT TAAGATGGGG GACATCCAAG CACCTGGAAC AAAAAGGACA CTAAGAATGG GAGAAGAATA CACAAAGGGA
GGTAGTACAG GGCCAATAAC AGATTITIGG AATTITICAA ATTICICTIT GAAGTAATTI TACAGTCAGT AAATGGAAGT
GGAAAAGAGG AATAGAAGAG CATTICATTG ATTITITITIT TCTCTTGTAC TTACACATCT CATGACCTCA TGTTCCCAGA
ACTTAACACT TAGTTGGGTT CTAGTAGGATA TTTTGGGTTG AAAAGATGTT TGCTGTTTTG CATTTTGTTC TGTTTTGTTG
GCTAGCCTGT GAATCTAGCA TTGTACGTGA GAAAGTGCAT TTCAGATTGA AAGCAACTGG GTTTTGGAAA TGAACTTCAA
TAACATATCC CAG

SEO ID NO:432: (Length of Sequence = 292 Nucleotides)

TTCACCGIGI TAGCCAGGAC GGICTOGAIC TOCTGACCIT GIGATCIGCC CACCTOGGCC TCCCAAAGIG CIGGIATTAC AGGCGIGAGC ACCGCGCCCG GCCACCATIC ACTAATTITC AAGAAATGIG GAAGIGITCT ATATTINCIT CCCACTCCAT AGCTCCAACA TIGITGGCIA TIATGAATTI GGCIATTAAG TGATGCCAAC AATATTTAAT GAAAAAAAGA TATAGCAGTA TAGTTGAAGG AGGAAGCTGA AAGAAAACGG TCCATCNGTG AGGAAAAAGGC CC

SEO ID NO:433: (Length of Sequence = 335 Nucleotides)

TTTTTTTCTC AGCAGAGGAT TTTATTGGTG GTCACCTGTG GCACAGGTTA GAGGAGCCGA AGTGCTGTNT TTGTGGTGGG
GGGGGGACCA CAAACCCCGG CCCTGCCCTC TTGCTTACAT AGGCTTCCCG CCTAGAAGCG CANCATGAAC ATGCCGCTAC
GGATCCGGTT GTAGTCTGGG AGCTGCTCAA TGGGGCCATA TCCAGCCACT GCTGGGGCAC TGGTCATAGA TGTACTTNGA
GCAGATCTCA CGTACCACAC TGGCATCCAC CTCCGCAAAT CCGGCTTTCC CATTCAGCCA GGGGGGNATG CGGGNGGGCC
ATAGGTCAGG AGGCT

SEO ID NO:434: (Length of Sequence = 390 Nucleotides)

GITGCIGACI GCIGATIGGA GATGACGIGI ACCCATCCIC TAGACAGICI GIGCITITCC TGICITIGGA GCITCCAGIT
CCACCCCCAT CAGITITITI CREACCACIC CATCITGCCI TATTICTCIC TCITTCCITI TGACTGGAAG AGIACICANC
TTTTCTAACA TCITTTCATA AACIGITTIG ATTICACITA TATTGATITI NAACGIATAA TGIGCIGGIG TICIATITCC
TCAGTTAGAT CAGAAGGCCC CTAAAGACAG GGCTCCATTG GIGTTAAACT GCCATCITCA AGGICTGGGA CITGATITCN

CTITITINAC CINCACAACA AGGCACTCCT CTTGCACCCA GTCGGAATTT CAGTGCCTGT GGGTCAAAGT

SEQ ID NO:435: (Length of Sequence = 427 Nucleotides)

TCACTAAACA GTAGATITAT TITATGTAGA TITGTITITC TATAAAAATA TATTTATGTG TICACAGGAA AAAAGITGAG
TTGGTATGTG GGGGTGACTT TCAGATACAT AATTAGITAA AGGITTGCIT ATGAAGITAG AAGGCATCIT AGCTTITATC
ATTITCCAAAT TITTCTTCAT AAAAAAGAAC ACCTGCCTAA TGTTTATGTA CAAAATCAGT GCTGGAGCCA GGAAGAGAAT
TTGGATTTTC CCAACCCTTG GACAGTTCTC TAGGGACTCA TGCCCACCAA CCATTCTTGA GACTATATAC AATCAATTAC
ATTAAAATGA TATTGACAGT AGACTAG

SEO ID NO:436: (Length of Sequence = 249 Nucleotides)

TCAAATAACC AGGAGGGGA CAGAAGATGA TGGCAAGGCA GACTGGGCAG TGTTTTNTAG ACACAGAACA AAGAATCAGA
ATTTGAAAAA AGANGAAAAA CAAATCINGG CAGCTGCAAC TTTAAAGTAT CACCTTTATA GATGGCAGG ATTTCCATTA
TGCAAATGGA ATCTAAGATT TCAATGTGNA ATCTTAGAAT GCAGTTTTAC CACTTGCAGT CINGTATTTG TGGTGGCCAT
GTGGTGAGT

SEO ID_NO:437: (Length of Sequence = 404 Nucleotides)

GICATTCACC CTAATCCCTC TITCACCTTC ACAGAACTIT CACACTCCAA TGTACTTGCT GITTGTAGAT GCTCCTATAA ACAGAAAGCT CTGGGAGACA GGTGTCTTGT TATTCTTGCT CTCTGTCATA TCTCTGGGCC TATCACAAGT ACTCAAAGCA TAGAAGTTCA ATAAATATGT GTTCAATGTA AGAAATGATC AGTGATTCTC AAGCTGCAGT GGCGTCAGGA TAACCTAGAC AGCCTGTTAG CACGGNTCAC TGNNNCCCAC CCCCACAGTT TCAGGTCTGG TCTGGGNTGG GGCCCAATAA TCTGTATTCC TAAAAGTCCC CAAGCAATGC TGGTGCTGTT CGTCCAGGGA CCATGCTTAA AGAACCACCC GGAATAGGAC TGGTGGACAA AAGG

SEO ID NO:438: (Length of Sequence = 337 Nucleotides)

CIGCAACITA TACCITCCAT TIACIAAAGI COCAGIATGI GICAAAGIAG TITICATICC TCACAGCCAT GITATGAGCI
AAATATCACI AACITICCCI TICAAAGGIG AAATAAACIG AGACICICGA AGATTAACII GCCCAAGGIC ACCIAGCICG
TIAGGAGGCA CAGGIGGGAC TIGAACCCAG TICTITCIGA ATTCAAAACC TCCAAAATGI CIGICACATC AAGCIGCITC
AATGAGATGC TAGAAAATCA GGACAGIGAG CAAGCIGGAG ATAANGGAAG ATATGGAGGA ACACGGGAAG TGTGATCCTC
ACACACATAC CCIGCAG

SEO ID NO:439: (Length of Sequence = 380 Nucleotides)

CATCGIGIAT GAAGGIAGCC ATTITGIACA TGITACCITG TIAAAAACAA AAGAGCAGCA ACATGITIAG AGIGGIGICT ATAGATAGAA CACTGCIGIT ATGITTAAGG AAAATTGGGG CGGGGCAGA AAAGATCAAT ATGACTAGIT AGAAGACTAT TAAGGAGAAC TITGIACATG AATTATGGAT GIAAGAATTA GAAAAAAAAA GATGATCATG TICAGAATTI TAGCITTITIT ACAATTGTAG TGGAAAAGAA AACTCCTAGA GIAATGAATC AATGGTATCC TACAAAAAGA GAGGIGCCAA AAATACCATG AAATATTATA TTAAAAAATT CACACGNATA GGTAGTTATA ATATGTAAAG GCCAGACTTC

SEO ID NO:440: (Length of Sequence = 335 Nucleotides)

CONTERGET THATTGACCA GIGGACTGIG ACTITIGATG TAATTTTATT TITGAGAGAG GGTOTGCTC TGTCACCCAG GCTGGAGTGC AATGGGGTGA TCTTGGCTCA CTGCAACCTC CGCCTCACGG GCTCCAGTGA TTCTCCTGCC TCAGCCTCCC GAGTAGCTGG GACTACAGGT GCACACCACC TTGGCTGGCT AGTTTATGTA ATTTTTTGTA TGTCTGTGGA GACAGGGTTT CACCATGITG COCAGGCTGG TCTCAAACTC CTGAACTCAG GTGATCTACC CGCCTTCCAA AGTACTGGGA TTACAGGCAT GAGCCACCAT AATAA

SEO ID NO:441: (Length of Sequence = 356 Nucleotides)

ACTAATIGIG TITCIGCITC AACCIGCATT TOCAGAGGIG CCTGITGGIC TGIAAITGGI TCIGGCATGI TIATAGGIAT TACAAAACCA AGICITATIT TGCATITCAC AGGATITAAG ATGAATAAAG TGATGIGGIT GIGCIAGGIT AGAGITGIAC AAACTIGACT CCCATCGCG ATGGIGGGIT CCCAGGCCTA CAACCTGACC TCIGCCCTCA CGCCCATCGI CACGCGCTCC CGGIGGCTCA ACGAGGAGCC CCTGACGCIG GCGGCCTTC AGCAGGCNCC CGGCCAACCT CAGTGACGIG GIGCAGCTCA TCTTTCIGGG TGGGACTCCC AATCCCCTTT CCCTTT

SEO ID NO:442: (Length of Sequence = 371 Nucleotides)

GATGATTITG TATCITITIC TATTIATIGA GATAATCAAA TGATTITIGI CCITCGITCT ATTGATGIGA TGITTATIGA
TCATGITTAT TGATTIGCAT ATGGTGAGCC ATCCITGIAT TCCTGGTATA AATGCCACCT GATCATGGIA TATNATCTIT
TINATGIGCT ATTGGATTIG GITTGCCAGI ATTTTGITGA GAATTITTTC ATCTGTGTCT ATTACGGATA TTGGCCTGIA
GTTTTTTTTG CTGTGTTCTT CTTTGGTTTT GATATCAGGA TAATGCTAGC TTTGTAGAAT GAGINAGGGA GGAGTTATCT
ACTCTTCAAT TTTTGGGAAC AGTTGCAGAA CTGTTGTGTGT TTTTAGAACA G

SEO ID NO:443: (Length of Sequence = 329 Nucleotides)

TGAACTGCCT TTATTTTTIN ATTICCCATC CAGAAACCCC AGTGTGATGG TGGAAGCAGC ATGAAAACAA CATCTCCCCA
GGCCTCGCAG TAGAGGCGAA GGGAACAGAG CTGCCCATGT GCCTGINTCT AAAGACGCCA CCCTCAGGTT GATGTCACCT
GTGGGAGACC GGGTCCACCT ACAGACACCA GGTGATGGTC CACCAGGCCC CAAGCTCCAG CCTGCTGAGT CCCCAAGACA
CAGGCTCATT AAATAGCTTC GTACAAAAAAC CCAAGGGTGT CCCTCCAGCT GGTAAAAAAT TGGGCAATTT CTACTTGGAG
GTCTGCTGT

SEO ID NO:444: (Length of Sequence = 358 Nucleotides)

TITITITITIA AGIACATAGG TCITIATITA AACACTGATI TITITITITAA ATATATACAC ACAAAACTTA GITCAGCAAG
GCITCATGAT ATACACCAAT TOCAAAATAA AACAATCAAA TGGICCAGGI GIAGAATGCC AGATTCCTIT TATCATCIGC
GAGGAAAAGA GAAGCAGGAT GAGGAAGAGI GAGGGAAGGC TCTGCCCAGA NGAGCTGCCG CCTCCTGGCA
CAGCAAACGC TCCAGGCCTG GGCCCTGTTC ATATCTGGAG TCGGAGGGAG ACTCCCATCG GCCGCTTTGG GACTGAAAGG
CCCCAAGGCTG TCACCAGGTC CCGGAAGAGA GGGAGGCA

SEQ ID NO:445: (Length of Sequence = 302 Nucleotides)

TCAGAACGGI GAGAAATAAA TIGCIGITGI TIATAAAGIA ACCIGITIAT GITATITITI TATAGAAGCC TGATCAGAAT

AAGACAATAT TGGATAGAAT ATTCAGGAAT GICITGCCIC CAATGITGGC CCCCCTGTAC TGAGCICTAA TCTACACTCA

CCTAAAAAAAT TATAAAATCA TAATAAAACT GAAAAAGICA AACTCICAAT TGCATCCCAG CACAAATATC ACAGNIGNIT

ATTTAAAAAAA TTATGTCAAG GCCCTAAAAA GCTAAAATCC NCAGNICTGC TAATATTTCT CT

SEO ID NO:446: (Length of Sequence = 367 Nucleotides)

TTACTICAGI AAAACAAGCC ACAGCAACAT TATGCTCTGC AGAGTCTTCT GTTCACCTTT GGGATGGAAA AGAGCTGCTT CTCCTAGGGN GGCAACTAAG GCCCAGGACC AAAACTCCCA TCTCCTA

SEO ID NO:447: (Length of Sequence = 295 Nucleotides)

CTGCAAACCC TTCAGCATTT AGCTAAAGTT ATTTCACAAT TCAATGCTTG TCTTGCACTG TCCTGGTCAT TTAAAAACTG
GTATCTCTTC AATAGCAAAT AGTATCATAA CAGACCACTA AATTTGGAGG GAAAGTGGTT TCTATTGCAG ATGGATGTAA
TTAAAAATTGG TGTAAATCAC AGGGTACAGA ATTCTTATCT GGTAAGAATT CTGACTTTTT TTTTAAAGAA GAAAAAAATAT
ATCCAGATCT GTATCCACAT GCTATTTAAA TGCTCAGGNC AAAAAGAAGC CACTA

SEO ID NO:448: (Length of Sequence = 233 Nucleotides)

CAGAATCAGC CCAAATGCCC ATCAATCAAC TGTGCATAAA GAAACTGTGA TATATATATA TCATAGAAGT TCAAACAGAA AAAATACAAA AAACTTAGCA GAGGATTGTA TCCTTTGCCG TTTATTTTGA TGACCATGCC ATCTTCTAAT CCCCAGAAAA AAACTGGAAA ACAGAATAAA TATAATTTNC TGATTATNCT TATGTAACAT AAATGGAATA TATATATATA TAT

SEO ID NO:449: (Length of Sequence = 341 Nucleotides)

ACTICCTICC TCAGGCTCCT GTACCAATCT TCAATTCACT TGGGATGTCC TAGTCTAAAA CATTTATTTC ATTTGAAAGG
AAAAATATCA ATTTCTATCT AAATTGGAGT AAGATCAAT TCAGATGTGT TTATTTACAA AACATAAGTT TGTTATTTAT
CTGTGTTTAA TTTGATCCNG GAACATTACA TGTAAAGAAC ATTCCATGTA AAGAACCAGG CAACTTGGCC AGGCATGGTG
GCTCACACCT GNTAACCCCA GCACTTTTGG GAGGGCCAAG GCAGGTGAAT TGGTTGAGAC CAGGRGGTTC AAGACCCAGC
CTGGGGCCAAA TATTGGCGAA A

SEO ID NO:450: (Length of Sequence = 313 Nucleotides)

TTTTTTTTT GACACAGITT CCAGTCCIGG AAACCTTTAG CTAATCTTTA GCATTCCITC AATGGIGGGA ATGGGCAACA
GATCACCATA GIATTAATAC TCIGIGIAAT TITATCACTA GAATGGITAA TTTCCATATC ATAGIAGAGC TGITGCAGAT
ATTTIGAAAT CCCATTATAC TCACTGCCAC TTCAAGATTA CIGIAGITGI TAGAACAGCT GCIAGATCIT ATTACITAAT
AAATTAATAA AGTGIGAATA TAACIATATA ACCATTITNA AAATGITTTT TGGATAACIT TCAATATAAT TGG

SEQ ID NO:451: (Length of Sequence = 351 Mucleotides)

GOGCCOGCTC CTGGGCACCC ACCCAGCTCA TTCGCCCGAGC GGCTCCCCTC CTGGGGTTGA GTGTCCTGGG CCTGAGTCTG
CAGCCTCAGC CATCTGTTCC CCAACTTGAT CTCCCACTGC TAGTTACAAA CAAATCGCCC GGCTTGTGCA AACCTCCTGG
GCTCAGTCCC CAGTCCCGCG GGGCATCATT TCATTCTTTC CTAGCCTGTA AGGTTTCTCC TGAAAAAATCT ATTGTTAGTC
TAATATGAAT TTCCTAATAT GTGACTTAAG GCTTTTCTCT TGCTGCTTTT AAAATTTTCT CTTTTGTCTT TGACTTTGAC
AATTTGGCTA TAATGTATGT TGGAGAGGAC C

SEO ID NO:452: (Length of Sequence = 363 Nucleotides)

GACAAGGAG AATTCTTGCT TTACCTATEG ACTGGCTTAA GCCGTGTGGC ATCCGAGGAA TGTTTCAAAT GTGTCTGTGT
TTCTCTTTAC ATTCCTTATT GTACCTCATT GTTCAATTCA CTTTTGTAAA TTCCACCTAA CATTTAATTA TTTTAAATTT
CTCCGTCATG AAGTTATTTT AAGACACTGG AATAAGTGCA GCTTTGTTTA TAACAGCATA GGATTATAAA CAACCTAAAG
AGTCAGCAGT GACATTGATG GCACATGCAT ACAATGGAAT ATTTTGTAGC TGTTAAAATA ATAANGAAGA TCCTGCTCTG
TGTATTTGAT ATGGGAAGGC CCCCCAAGGT CTACAGTTAA GGG

SEO ID NO:453: (Length of Sequence = 382 Nucleotides)

ATGAGGGAAA AGATGGTGCC ATTGAAGATA TTATCACAGT GCTGAAGACT GINCCCTTTA CTGCTCGCAC CGCCAAGCGT
GGCTCTCGGT TTTNCTGCGA ACCTGTTCTC ACTGAGGAAT ACCATTACTA AACTATTACT CTTTCTCACC TGATGCTCTT
AAAAGATCTT AGAAACCAAC CATACAGACG AGCCGATGCG GTGAGGAGAA GCGTCAGGCG GCGCTTTGAT GATCAGAACT
TGCGTTCTGT TAATGGTGCC GAAATAACAA TGTGAACCTG AGACTGGCCT GCAATGAATA CAGGGTGTG CGTGTTCAGG
AGGTTTTCTG TTGCGGTCAC CCATGATGGC GGGCCINCCC ATTTGGGCCA ACTTTTCCTG GG

SEO ID NO:454: (Length of Sequence = 391 Nucleotides)

SEQ ID NO:455: (Length of Sequence = 282 Nucleotides)

TIGAGIACTC ATTIGAGGAC TGCAGTCATA GATTTAAAGT GIAATCAGTC AACTCAGTGG AATTACTTTC TCCATTAATC
TTAAATTGCT TCAGGACTGT TTCAGCCTAA GCCAGTAGCT GGGTTTAACC AAATTTGAAG ATTITINCTAG GAGAGTTTGG
CACGAGGAGA GAGGGGCAAA GGCGTGTAAG GCAGTGTTTA TAACAGTGGC CCATGGAATT GATCATGGGT AAAGAGAAAA
CAAGGACATG CGAGGAGGTG ATAAATAGAN CAAAACAAAG CA

SEO ID NO:456: (Length of Sequence = 340 Nucleotides)

CTAACTTATG TTTGAGATCT TCAATGAAAT TAGTTACTAA TATTINGCTT TATTCTTCTC AAAAGATTTA ACATGATAAT TCTGACCTAA TCCAAAAAAA AAAAATTCAT GGGCCACTGT TTTGCATGTA ATATGTAAGA NCTCACCTTG ATGTTAAACT CCCAACCCTTG GCTGAAACAG GTTAATGATC ATTTGTNGTT ATTTATTTCT ATAAATAGTT TGAAGTTGGC CAGGCCTGGT GGCGTCTCGC TGTGTCTCCC AGGGTTGGAG TTCGGTGGGG CAAATCTCGG CTTCACTGCA AGCTTCCGCC TCCCCGGGGT TCACACCATT CTTCCTGCCT

SEO ID NO:457: (Length of Sequence = 338 Nucleotides)

ATGAAAAAGT CTCCAGAGAT TATCAGTGGG CGGATGACAT TTGCCCTCTG TTGCTATTCT TTGACATTCA TGAGATTTGC CTACAAGGTA CAGCCTCGGA ACTGGCTTCT GTTTGCATGC CACGCAGCAA ATGAAGTAGC CCAGCTCATC CAGGGAGGGC GGCTTATCAA ACACGAGGATG ACTAAAACGG CATCTGCATA ACAATGGAAA AGGAAGAACA AGGTCTTGAA GGGACAGCAT TGCCAGCTGC TGCTGAGTCA CAGATTTCAT TATAAATAGC CTCCCTAAGG AAAATACACT GAATGCTATT TTTTACTAAA CCATTCTATT TTTTATAGG

SEQ ID NO:458: (Length of Sequence = 370 Nucleotides)

GITTICITIC GGAGCIGAAC CAAAGAATGI GCACCCICII TCICTAGIGC TGIGGIGICT GCITATITIT GIATTIGIGC
TTICCATCCA TCITCIGIGA TCACAAGGCA TTCITAAGGI TTICTAGCAC GACTIGCGGA CATCCAGACI CGIGGGGGGC
CCACCCATGG CTCGGTAAGC CAGCAGCCCA GGGCACTGGC ACTACCATGA GGCACTGCAT TAATTGCTGC ATACAGCTGI
TACCCGACGG CGCACACAAG CAGCAGGTUA ACTGCCAAGG GGGCCCCCAT CACCGTCACC AGGCGTGCCC CACGTTGCAA
AGGAGGAAAA ACAAAATTCC TGGTTTCCGT GTGGGACAGI AAAGCAGATG

.

SEO ID NO:459: (Length of Sequence = 339 Nucleotides)

ATTITCCTAG AACTGAAATC ATCTACGGIT CTCAGAGCTA AACTTCCAAA GCTACAGTCA GCAATTITTC ATCAGAGCCC
AAGGGAGAGG GGCCAGGGTA AAAGAGACGA GACTGTAGAG AGGCATAGAG AGACCAGTAG GAAGAGGGTG GGAGAGGGCA
CTTATTTCTC TCTGTCCTCT CAGTGGGTTA CAAATCAGAT CTGGTGACAA CACTGAGGGG GCCAGGTCAG GGTATGTNGA
TGAGAAATGA CACTGGAAGG AACATCAAAG CCGCAGCTAC AAAAAGAAAG TCATCAAGCC CCAAATAGAA GGGGGAGCCT
CCCAGTGCAC CTCAGAAAT

SEO ID NO:460: (Length of Sequence = 380 Nucleotides)

GAGCITTIGC ACTGCAAAAG GAACAGTCAG CAGAATAAAC AGACAGTTAG AAGTACITCC CTATGTAGAG ACACACTCAA
GTGAAAGGGA ACCAGGCTCT ACCACTTGAA ATAAGGAGTA TCAAGGAACT TGTGGACAGC TTTTAAAACT ACCACTGGCA
ACTAGGTCTT GAGGTGGATA AATGAAGAAA TTTGGGGAAT CTCACACTGG AGATGTTTGA TGTAGGTAAA TGANCTGAGA
TTCATTAGGT GTGAAATAAT GAAGTGTATA TATAGTTCTG CATATACATG CCTGGGGAAG GTATAATATT CAGAGGCATA
CTATCACTCA ATTTGTATCT GCTGTGGGCC TCAGACAGTA CAGGGGCAGT GTTTGCATTG

SEO ID NO:461: (Length of Sequence = 317 Nucleotides)

GTCATTAGGA AGCCTTTATT GGGTTATATT CAATTTGACC TCCCACCAAA TTAAGCGGGA AAAAACAAAA AAATAAGAAA TCCCAGTAAA AGAGCCCCTC AAGATTTCAT AAACTACAAA CTAAAGCTGC TAGTTAATAA GGAAATGGCA GAATTTCAG AGCTGTATAA TACAAAAAATT CCTGTAATTT AAGCAGATGT TTTCCTCACT GATGACAAAT CTTCCAACAC AATGTGAAGT TATGCTACTT GGGATATTTG TAGGCAAAAC CATTTTTTTT TTGTACAAAA ACAAAAGCAA GGGACCNTGG AAAAAAA

SEO ID NO:462: (Length of Sequence = 261 Nucleotides)

AAAAAGGCCA TAAATCCTIN CCCTCGTGGA GCTTACCTTC TAATAAGGAG AGACAGAGGG TNAGAAACAA ACAAACAAAA ATATGTNAGT TAACACAGAG TGTTGGAGGG TGTCAGGTGC TATGGGAGGA ACGTGGAGCA TGTCAAGGNG AGAGCAGGCA AGAGGGCATT CTGGAAAGGC CTAGGANGAT GGTGACATTT TACCTTCATA TCCACCAACC CCCAGCACAA AGCATTTTCC AGAGGNAGNC AGAGGAGGGC A

SEO ID NO:463: (Length of Sequence = 387 Nucleotides)

ATACAAGTAC ATCCAGGAGC TATGGAGAAA GAAGCAGTCT GATGTCATGC GCTTTCTTCT GAGGGTCCGC TGCTGGCAGT ACCGCCAGCT CTCTGCTCTC CACAGGGCTC CCCGCCCCAC CCGGCCTGAT AAAGCGCGCC GACTGGCCTA CAAGGCCAAG CAAGGTTACG TTATATATAG GATTCGTGTT CGCCGTGGTG GCCGAAAAACG CCCAGTTCCT AAGGGTGCAA CTTACGGCAA GCCTGTCCAT CATGGTGTTA ACCAGCTAAA GTTTGCTCGA AGCCTTCAGT CCGTTGCAGA GGAGCGAGCT GGACCACACT GTGGGGCTCT TGAGAGTCCT GAATTCTTAC TNGGGTTTGG TGAAGATTTC ACATACAAAT TTTTTGA

SEO ID NO:464: (Length of Sequence = 397 Nucleotides)

AGGCAGGAGG CCGGGGAAGT GGGGTGGGGT CGCGACACCG ACAGCAGCT CAGGCAGGC CATGCTGCGC TCAGCTCCCT CAGGCTGTCA CTCTTAATCA TCATGTCACT ATCTCTGGGG CGTGTCAGTC ACCATCAACG ACGTGTCCC CAAGCTGCAG AGGACGCAAA TCCAGCTCTC CAAGAGGCTC TGTTGGCCCT CTCCACATGG GCTTNAGGGT CAAGGGTTGG GGGCACGTTC GGACCGNCCT TCCTGNCTCT TTNGAAGAAG ATCCTCCAAN GTNCCCGGCT TCAGCTTCTT CCGGGCCTCT TTTTGGCA

SEO ID NO:466: (Length of Sequence = 352 Nucleotides)

CATTGIATTI CCCTTCTICA AATTAATAC CIACCAAAAA ATGGAAAAGA ATTTTACATG CACTITAAAA TAGIAAAATG
GAAAGIGAAT TITTAAAATA TATGCATTAA AAGITTACTT TAATTTCCAG TGGGACTTCC TITATGAAAT TITCCATAAC
CTCTTCCTGG AGTATTACAA GATCTCCAAC ATCTCATAAA CTAATTGIGA TATTAGIGGA ACCATAAGCA AATGTATATT
TTTAGTGGAA ATAGATTATG AATGAAAGCC AAGCACCTTA CTTTAAAGCC AAAATATGAG ATTTTCCATT AAAAACCATT
GGTCCATAAT AGGGAGGGGG GTTTTTTTAAT TT

SEO ID NO:467: (Length of Sequence = 352 Nucleotides)

TGAAAGGCAA AAAATAAATA AATAAAAATA AATACCATTT GCAGAGACAG AGAAACCATC AGAAGAAGAC AAGCAAGGTT
GTITGAATTA CTACGCCTAG AATTTAGAAT AACTACTATG ATTAAAACGA AAAAGGCTTT AATGGATAAA ATAGATAGCT
CCTAATAACA GATAGTAATA ACATATGGGT AATGTGAGCA GAGAGATGGA AATCTTAGAN CAACAACAGC AACAACGNCA
AAGCGTTAGG GATCAAAAAAC ACTGTAACAA AAATTAAGAN TCCCTTTTAT GGGCTINITA ATAGNCINGG ATACAGGTAA
GTAAAGAATC CCTGTGCTTT AAGGAGCCAT CA

SEO ID NO:468: (Length of Sequence = 336 Nucleotides)

TGACATCIGC ATCTIACATT ATTAAATGCA AAGGAATATC AAAGACTCCT CIGCIAGAAC CATTITIATT CATAAAGTCA
CATTATCATT GIAGAAGTCT TGIAAAAATG CTACCIGAAA TGAATTATGT CCGICTICCC ATCIGGCITA CAAAATTCTT
GAGGAAGCAT CIGCCTCGTA GCTCTTTATC TTTCTATTTC CTACTACAGG GACAATGTAT ATGGAAAGAT AAATGTGTGT
AGGIGTATAA ATTCTCAATA AATATTIGCT GAATTAGATT GIACAGTTGT TATCTTTTAA GNITAACTCA TCCTGAGGTA
CATTITATTA TTGGGC

SEO ID NO:469: (Length of Sequence = 156 Nucleotides)

GACCGATGIA GAATICIGIC TGGAGACGIT CICCCCTICA ATTCAATGGG AAGGNICIIT TCIGGCATGA NCICICCGAT GTCIAATGAG CICTGAGCAC CATCCATAAG CITINNCACA TTCITIANAT ATAAAAGGIT TCICICCACT GIGAAT

SEO ID NO:470: (Length of Sequence = 350 Nucleotides)

TTCTCATGTC TGAATTTCAC ACGCACAAGT CTGAAATGTG AAGGTTTCTT AATGTTGGTT TTATGGTTCG TGTAAGATTT
TTGGGAAATG AAGGCTCTT CATTAGGATA AAATGGTCTT AACTTCCCAG AGAAGAATTT CCTGACAACG TGGCTGAAGT
TAGATACAAA TGTTAATATA GAAGANTGCT TTTATTTGAA TTTCTAGCAA ATGGTTTTCA ACTACTTTAA ATATGACCNA
CTTGAAAAGTA TTATTCCINT TTTAAAACTA CTTTTNATGT ATAGATCTAA GGTCTGCTTG AAGCTAGTAG GTTAAAGTGT
TTGAGAAAATA AAGGCAAGAT INFINCNITIA

SEQ ID NO:471: (Length of Sequence = 270 Nucleotides)

GGAGCAGGC TGGGAGTCAG TGGGAGATTG GGAGTCCAAG TCTGGACATG TTACATATGC TATGTCTATT ACAGATCTGA GTATAAATGT GAAGTGGAGT TTTACCACGT GATTCTGAAG TTCAGAGAAG AGGTACAGAT TAGAGATAAA GATTTNGGAG TCACAAATAT AAAGATGTAT GACTTNATGA GATTACCAAG GAAGTGGAGA TTAATAGCAA AAAGAAAAGT TTCAAGCTTC AAGCCCCGAA GCATTCTAAT GTTTACAGCT SEO ID NO:473: (Length of Sequence = 345 Nucleotides)

TITATIGIAG TICAAATACA TAAACIGAAC ATICAAACAT CITAAAATTA AACITTAGCA ACAAAGITTA ACATICAAAC
AGGAGIATAG TITACAAGAA ACACCCAGAA AGGIAATTIG TIGICTAATC CAGAATATIG ATAAAGATCA CITAATGGIG
AATAAAATAT GITTAACCAG TGGITCTATT CIGGCCAACA TGITAGITAT GACCGIGGIT CCATACCTGA GAAGAAATTA
CTACATAAAT CITCICITAG GCTAAACAAC ANGACTCGGI CIATAATTCA GAGGGGNTAA TCAAAGCACG TAAGGGIACC
AAAATAAAAC TAATCIGATC TITAG

SEO ID NO:474: (Length of Sequence = 433 Nucleotides)

CAGAATTAGA GCIGIACCCC AAGGGGGAAT TCTGTCCTAG GAGACAGTGA GINCTAAGTA CACTCTGGAC AAGCACCAGA
CACAGAAGCT GCCTCAGTTT GIGCTCCCCC TGCAAAGCAG AGCCTGAGAC AAGGATTTGG GTACAAGGAG TTTCACTCAA
TATTATATTT CCAAGATGCA CCCATGCTTT ATATGGCTAT AGTGCATCCA TTTTACTGCT TTATACTTTC CATTAGGTGA
CTATATTAGT ATATATTTAT AATTCCTAGG TCTTTTTGTT CTCTTATTTG TTAATAATTA TAAACTCCAA GCCCATTGTG
GTAGATTGCT ATTTCTCAGA GATATTTTCT GCTCCTTCCT GGGGGACAAT AATACTNTTC TCCCATCAAT GGCAGATGIN
GGGCTTGTNA CATTTTCTGG TCAATGGAAT GAG

SEQ ID NO:475: (Length of Sequence = 427 Nucleotides)

GATATEGITI GIGIECCCAC CCAAATCICA TCTAGAACIG TAGITICCAT AATCCCCACG TCGTGGANEG GACCIGGIGG
GAGGIAATCG AACCATGGGG GIGGITACCI CCATGCIGIC CITATGATGG TGAGITCICA TGAGATCIGA TGGITITIATA
AGGGACITTI CCCCCCTTIG CICTGCACIT TICCATGCIG CCACCACGIG AAGAAGGATG TGITIGCTIC TCCTTCCACC
ATGATTIAAG TITTCINAGG CCTCTCCAGC CATGCIGAAC TGIGAGICAA TTAAACCICT TTCCTTTAAA AATTACCCAG
TCCCAGGNAT GICTTCATTA GCAACCICAG AGCAGATTAG NCACAATTCC ACAACTTGGA GAATNGGIGT TCAAGITTCA
CTCTGGCCIT NAACAACCCA AAATTTA

SEQ ID NO:476: (Length of Sequence = 351 Nucleotides)

CGCCGCTAGG GCGCCNGGGG GTCGGGACGC CGGGCTAGGG GCGCGTCATG TGGCCGCTCA CGGTCCCGCC GNCGCTGCTG
CTGCTGCTGT GCTCAGGCCT GGCCGGACAG ACTCTCTTCC AGAACCCAGA AGAGGGCTGG CAGCTGTACA CCTCAGGCCA
GGCCCCTNAC GGGAAATGCA TCTNCACGGC CGTNATCCCA GCGCAGAGTA CCTGCTCTCG AGATGGCAGG AGTCGGGAGC
TGCGGCAACT NATGGAGAAG GTNCAGAACG TCTCCCAGTC CATGGAGGTC CTTNAGTTNC GGACGTATCG CGACCTCCAG
TATGTACGCG GCATGGAGAC CCTCATTCGG A

<u>SEO ID NO:477:</u> (Length of Sequence = 333 Nucleotides)

GGTCTCACTC CGTCATCCAA GCTGGAGTGC AGTGGTGCAA TCCTCAACTC ACTGCAACCT CCGCTCCCGG TTTGAGTGAT
TCTCATGCCT CAGCCTCCCG AGTAGCTGGG ATTACAGGCA TGAGCCACTG TGCCCAGCTG GGATATAGAA TCTAAGAGTT
GATTGTGGAA AACACGTGAA TCTATTGCGC GCATTTNTCA TTTAGCAAGA TGGCAGCAGT CCAGCTGTTC TTTGCAGCTG
GAGATGAACT TTTAAAAATC CCCTTCACAC TTAATGTACT GACCGAGACA GAAGTACCTG AAAAACAGCT NTGCATGGCA
GGCCCGGCAA TAG

SEQ ID NO:478: (Length of Sequence = 458 Nucleotides)

ACATGITAAA ATAAGGIAAT ATGAAATAAT CIAAAAAAAA AAAAAGIGCA GAACCAAGAC CICTGTGATA ATCCTATITA
AAAAAATAGC TACAATTITA GITAGAATGT TICCCTTATG AGAAAGCATT TICTGCATAA CITTTAATGT ACTGACCTTT
TCCCAAGCTTG CIGAGCTGGC CITTGTCTCA ACTCACTTGG GACACCTTTC CCTGTGCCTC ACCAGGGCCC ACCCCAAGTC
CCAGTTTCTC TAGGGGGTCT CTCGGGACCC CITGAATCCC TTTNCTGATT TGTGCTGCCT TTAGCAGNCG GAATGGGCTG

GCAGACCACC CTACATNOTO CTGTGTGTGG GGACACTGTC AGGNTGTCCT CCCTGCATTA GNOTOTGCTG AGTTTCCTAC CATGTGNCCA GGATGGNGTC CATAGTCGGG GCATNAAGGA CTTAGGATGG GCCCAGTC

SEO ID NO:479: (Length of Sequence = 360 Nucleotides)

GCATCGIATC INCITIAAGA AAAACACITC ITCAAAATCC TACACIATGA AAAACIGICI TCAGGAATIG ITHATITGGI CCGITGATCI AGTGAGGCIG AGITCITAAA ICITICACCC CCAAGITAAA AATTGGAGCA ACAAAACAAA ACTCCAGCAA GGCATAAATA AGATATTAAA GIGCATATAT ACAATACCAG AAAAGITTAG ATTGGGAACA GCAAAAATTI CTAGIGCAAA AACIGCTITT GCCAGCAAAG CTCCCTCTCT GGAATCAAAG GGCTACAGTA AAAGITTAAAA TTGGAACAGG MITAAGCAAT GTCIGICTIT AGTCACAAGI NAATATATGT GCCATGCACCC

SEO ID NO:480: (Length of Sequence = 322 Nucleotides)

SEO ID NO:481: (Length of Sequence = 369 Nucleotides)

CCTGGGCAAA GCATTGATCT GGTAGCCTTG CTCCAGAAGC CTGTTCCTCA CAGTCAAGCC TCAGAAGCCA ACTCCTTTGA
AACTTCCCAA CAGCAGGGCT TTGGCCAAGC CCTTGINTTC ACAAATTCGC AACACAACAA TCAGATGGCA CCAGGGACTG
GCAGCTCCAC TGCCGTCAAC TCCTGTTCTC CTCAGAGCCT GTCATCCGTC CTTGGCTCAG GATTTGGAGA GCTTGCACCA
CCAAAAATTGG CAAACATCAC CAGCTCCCAG ATTTTGGACC AGTTGAAAGC TCCGAGTTTG GGNCAGTTTT ANCACCANCC
CAAGTACACA GCAGAATAGG TACAAGTCAA CCCTACAACT ACTACTTCT

SEO ID NO:482: (Length of Sequence = 255 Nucleotides)

GAGAGAATCT CECTCIGICG CCCAGGCIGG AGIGCAGIGG CGCAATCCCG GCTCACIGCA ACCICCGCCT CCCGGGITCA
AGIGATICIN CIGCCICGGC CICCCCAGIA GITGGGATTA CGGGIGCACA CCACCGCACC CGGCIGATIT TITGIATITIT
TGGIAGAGAT GGAGITTCAC CATGGCIGGG CIGGICTIGA ACICCIGATC TCAGGIGATC TGCCCGCCTC AGGCTACCAG
AGINCIGGGG TIACA

SEO ID NO:483: (Length of Sequence = 353 Nucleotides)

CTGGATAATC AGGCCATGT GCTTTAACAG GATGTAAAGG GGAAGCTCAT GATTAAACAT GGGAAATATG CAGCAAATTG CAAGACCTGA GCTTAACCGC ATAATTAGAA CATAATTTIN CACTTCTTCC AGAGCATCAG CCAAGCAAAG GACTGAGAAA TCTGCCAACCC AATTGTCCTA AAAAGAAACT TAGGCTTCAC ATTTGTGACA TAATTTCTTT TAAAATGAAT ATAAAATTTT ATTTTTNATA TTTGTAGAGC ATAGGATGAT TGAAATCCAG TTGTTGTTTT ATCTGACCTC CATATCTAAT ATGGCTAGTG CCGTTACTAC TCTACAGAAC GCGCAATAAG TCA

SEO ID NO:484: (Length of Sequence = 371 Nucleotides)

GACCCAGAAA ATGGAGCTAG CTACATTICT CACACTTACT GICATAATTA CATGITTATA TICTATTAGT TGIAATTATT
TITCACCTAT CCTCTCATTA GAATGITATA CCTATAGAGC AGATACCATT CCAGITTIAA TITTITIGCCC CGACTCCTAG
TAAGIACGIG ACCTATTACA GGGAACTTAA AACAAACAAA AAGTCTGCTG AGTCTGGGAT GITTTAAGGA TCGAAGGAAC
ATGITGGTCC AATTTGCCTT CACAGAGGGT TACCTCTGCT TITCTACCGA ATGIGGGAATT GCTCCCATGT GGATTTINAA
GGAATTCCAG TCTACCCTCA GGGGAAGGCC CACATGTAAT GCCAGAGGTC T

188

SEO ID NO:485: (Length of Sequence = 376 Nucleotides)

GGTCCGACGC TGTGTCAAGC TCTGCACCGG CCATGAGTAT GCAGCCAAGA TCATCAACAC CAAGAAGCTG TCAGCCAGAG
ATCACCAGAA GCTGGAGAGA GAGGCTCGGN TCTGCCGCCT TCTGAAGCAT TCCAACATCG TGCGTCTCCA CGACAGCATC
TCCGAGGAGG GCTTCCACTA CCTGGTCTTC GATCTGGTCA CTGGTGGGGA GCTCTTTGAA GACATTGTGG CGAGAGAGTA
CTACAGCGAG GCTGATGCCA GTCACTGTAT CCAGCAGATC CTGGGAGGCC GTTCTCCATT GTAACCAAAT GGGGGTCGTC
CACAGAGACC TCAAGCCGGA GAACCTGCTT CTMGCCAGCA AAGINCAAAG GGGCTT

SEO ID NO:486: (Length of Sequence = 396 Nucleotides)

TIGATATITG TGICTAATTC CAGCIACITI GAAAGCTAAG GCAAGGGGAT TACTGTATTA ATAAATTCTC ATGCTGITAA
TAAAGACATA ACCAAGACTG GATAATTCAT AATGAAAAAG GITAATGGCC TCACAGTITC ACATGCTGG GGAGGTCTCA
CAATTATTGG AGCAAACAAG AGACTTTGIT CAGGGGAATC TCCACTTATA AAACCATCAG ATCACGTGAG ACTTTTTTGC
TATCATGAGA ACAGCATGGG AAAATCCCAC CCCCATGATT CAATTACCTC CCACAGGGTC CCTCCCAGGG ACATGTGGAG
ATTATTACAA TTCAAGATGA GATTTGGTTG GGGACAGAGA GGCCAAACCA TATCAATTAC TTAAGGCTAG GGGTTT

<u>SEO ID NO:487:</u> (Length of Sequence = 375 Nucleotides)

TGATTAAAAT AATAGAGITT AGIAATATGG ATGAATATAA GATAAATATT TAAAAAGCAG TIGIATTITT ATAGCCCAGC AAGATAAAAGT TCAAATATGT ATTITITATA AAGATGGATT TACAATAACA TCAAAAATTA AAATGCACCT TGAAATAATA AAGACATGIA AACCCITITA TGAAGACAGA TTITITTAAAG CATTITITAAA AATACTITITT CATTGACAAA TAATTATCCN TATTINTGGG GIACACAGTA ATGITTCAAT ACATATAATA AATAGTGATC AGATCAGAAT AATCAGCTTA TCCATCATTT CAAACACTTA TCATTICINT GIGITAGGGG CCATTCAACA TCCTGCTTCT GGCTA

SEO ID NO:488: (Length of Sequence = 323 Nucleotides)

CACTGCATTA ATGATTGGNT TAACAGTATA TAAACAAGGG CCATGGTTT TTTTACTAAA GTAGGTCTGA AAGATCAATA
TAAATACTAA TGGGGGCAGG GAGGAGTGTT TTATACCCCA AACTCCAATA TTCCAGCTCT GTGTCCTGTC CTATTATTAT
AATTTGTAAA AATCTTAACG ACGCAGTGAT TCGAGTTTTC GTAACTTCAA TGATGTGTTA GAGGACAATG CATCTTGGTT
TGAAGAATTT GCTGTATCCG AAGGCCGGAA AAGTACTCGA CCACGATGAT TAAATACATA AAAGGATGGG TGATTCCTTA
CCG

SEO ID NO:489: (Length of Sequence = 326 Nucleotides)

TTACCITTTA CICIGATCAT AATCTCCCAC CIGICIAAGA GGITATTAT TCCTTATITA GAGGGCCICI ATTGCCATGT
GCCTGGAATT ATTATATGCT CATCACTTTA TGAAGAATAA AATTIGICIT TCCTGCITTA AAGITACATT CGITCITCCG
CTCAAATCCT GATCIGGICC ATTAAAGAGI GITCGCAGAC AAAGITTCIG AAAGATTAGA GAAGAATCCC CCCCAAGATT
GCCCCAACAC TGAACTACAG ACAAACACTA TTITATITTAA ATTAAGGNGAC AGCTTTCTAA AAGITATACAT TCCTCTAATA
AAAATA

SEO ID NO:490: (Length of Sequence = 186 Nucleotides)

CTCAGATCCA TCAAGATGTG AAACTCGCAA GITTGGTGCA GAGAAGGTAC ATGGGTTTCC TTCTTTTCTC ATCTGTATTC CCTTTTCTGC AATTATTTTC TTTGCCACAT ACTAGCCAGC AAACCAGGCA CCTTTGCCAG AGCCATTAAG CTACAAAAAT ACTTAATATT TTAATTTGAA CTCTGC

SEO ID NO:491: (Length of Sequence = 347 Nucleotides)

CCTGTACTTG TCGTCCCTCA TTCACTTAAT TATGATACTT GCCTGGCATC TTGCAGGTTT CTGATGCTGT TACCCCAGTA
TAGACCAAGT GCAGACAGAA TITCATTTCT GCTTTATTAA GGCACAGTCT TGAGAAACCC ATTGGCTTCA CACACAATTA
ATTAATTTINT GGCAACAAGC TACTATATTG GCTTGCATGT CACTTTCACC TCTCTGGGCA TTAGTTTNCT CTAATATTTA
TAAAAGAAGG ACATGACTTT CTAAGGTTCC TTGCAGTAAT TATGCAGTTC TATTCTAATA GATGCTTAAG CATAAAACCC
ATTTTAATAC TGTCCCAAGG ATCCAGG

SEO ID NO:492: (Length of Sequence = 320 Nucleotides)

GAATTIGENT CCAAAGITIG GACATIGCAT TICATTAATA CGICCCITAA GITTATITIA ATCIGIATIT TCCICCTCCC
THITIGIGITC TITIGIAATCI CITTIGCIG TIGITITCGG TIAAAGAAAC CATGITITIT TCGICCIGIG AGIGCTCCT
GITCAGAATT TIACIGATTI CATCIGCIGG TATCATITAG CATGITGCIC TGICCGCCGI AGIACITTAA ACIAGACGIT
AGATCIAGAG ATGIGATCIA CITCGGIAGG ACTITIGICAA GAATACTIGI AAGIAGGIAT TIACGIACCA GGGCNCACAT

SEO ID NO:493: (Length of Sequence = 339 Nucleotides)

TECCAAGTIT GCTGGAACAT TATCAGATGG CTTAGGGAAG ACGATGGACA ATCGGCATCA GTCAGAGCGG GAGTACATCA
GGTACCATGC AGCCACAAGT GGTGAACACC TTGTAGCCGG CATCCATGGC CTGGCTCATG GTATCATTGG TGGACTGACC
AGTGTTATAA CTTCGACAGT GGAAGGTGTG AAAACAGAAG GGGGTGTCAG CGGTTTCATA TCTGGCCTTG GAAAAGGGCT
TGTTGGCACT GTAACCAAGC CANTGGCAGG CGCCCTGGAT TTTGCATCAG AAACAGNCCA GGCGGTGAGA GACACAGNCA
CACTTCAGCG GCCCCAGGN

SEO ID NO:494: (Length of Sequence = 366 Nucleotides)

GEAGGCCITT GGAAAGTAAT TAGGATTAGA TAAAATCATC AGGGTGGGGC CACCATAATG GGGCTGGTGG CTTTATAAGA GGAAGAGAGA CCTGAGCTGA CACGCATGIN CTINCCCTCT TGCTATGTGG TGCCCTCAGC CATGTTAGGG CACAGCAAGA AGGCCCTCAC CAGATATTGG GGTGGTCTIN GACCTCCCAC CCTCCAGAAC TGTAAGAAAT AGATTTTTTT ATATATTACC CAGTCTATGA TATTCTGTTA COGNAACAGN AAACAGACTA AGACAAGCTT CTTAAACAAA TTGANAATAG AGTTTTAAGA TNCAGACTTT CATTGCCTTT AACAGGGGCC AAGAATATCT ATTTCA

SEO ID NO:495: (Length of Sequence = 384 Nucleotides)

CEAGGAAGGC AAGAAGCGCA GGGGGTGGCC CGCNTGGCGT CGGTGGCCTC CGCTCCTGCT CGCAGCCCCT GTGGTCAGAG CTGGATACAA GATTCAAGAC CCTTCTNTTG CTTGTNACCC GCTCCAGGTT GGAGCCACAG ACACCCACCG CCACCCCGGC TGGGTCTGCN TCCTTTCCTG TGCCTTTCCC TCCAGAATGC GGCCTCAGAC CTAGAAGCTC AACCCCCCTA TGAGGGCCAC GTCCTGGGGT AGCTCCTGAC CTNCGACCTT ATGTCCAAAT TTCACACCCA TGGTTTTTCA TTTGACCCGG CCCCTTCTCG CTCATAATGA CAACNAGCTT CCTTTGAGAG GGATCAGAGN CCAATTGCAC AAGGAGGAGC CGCT

SEO ID NO:496: (Length of Sequence = 342 Nucleotides)

TACCITAGIA AATGCAATIT TOGAACAGGC COCCATCITC AACIGGIATA GCATCITCCA CACCCIGIAG CCITCAAACA
TCACCIGITA AAATACIGCC CATTCCATGI CATGIATATC TGCCCATTTA TGGGAGCAGI GAGIGGAACC CIGACAGIGA
CGGACTITAA GCTGIACITC AAAAATGICG AGAGGGACCC GCATTTTATC CITGATGITC CCCTTGGAGI GATCAGCAGA
GTGGAGAAGA TTGTGACCAC AGAGCCATGG AGACAATTCC TGTGGTATAG AGATAGTGTG CAAGGATATG AGGAACTTGC
GGCTTGCTTA TAAAACAGGA AG

GATTTATTAA GTATCCCCGA AAATATAAAC ACAAACCAGT AAAAAACAAA ACCGTAAAAC GTCAGGCCTG GAGCTGCAAT
AAGACAGAGA CAGGAGCAGC TCACACGTGG CCTAGGTGGG GAGGACGAGG CCATAAATAC TGCAGGAGGG CGCCAAGGGA
GCCCTAGGGC GAGGGGAAAG CAGGGTGTCG GCAGCGAGAT GGNTCCNGGG GTTTAGACAC TGCTGGCTTC GGNCCCGGCC
GGCACCANGA CTCTCACTTC CAGCTGCGAG CAG

SEO ID NO:498: (Length of Sequence = 319 Nucleotides)

ATTCCCAAAA ATAGAGICIG GACCICITAC CGCTACAAAT TCCAGGITCT CAGGIACAGC CIGAGAGIAT GCAGATATAA
TACACCACAG ATGATICICT CCCITITITG TITTITITIT TITTIGAGACA GAATCICATT CIGICACCCA
GGNIGGAGIG CAGIGGGCIG ATCICGGCIC ANTACTICIC CCGCCTCCNG GNITCAAGCA ATTCICCTGC CINAGCCTCC
CGAGIAGCIG GGNCIACAGG NGCACACCAC CATGCCCATC CAATTITITGG ATTITAAGIA TAGITGGGGT TICACCATT

SEO ID NO:499: (Length of Sequence = 408 Nucleotides)

GAGAAATACC TAATGIGAAT GACGAGTIGA TGGGIGCAGC ACACCAACAT GGCACATGIA TACCTATGIA ACAAACCIGC ACATTGIGAC ATGIACTCIA GAACITAAAG TATAATAATA AAAAAAGAGA ACCTITAAAA AAAAATAGAC TGCCAGATAG ACTAATAAAT AAAAAAGAGA GGITGAAATA ATCATAAATG ACTAAGAGGA TGITACCCCA CAGAACTACA AAAAACAAAC AAAAAAAAACCT CAGAGCTAC TAAAACCACC CIATGCACAC AAACTAGAAA ACCTAGAAGA AATGGGIAAA TTICTGGAAA CATACANCCA CCGAAGATTG AACCAGGGAG AGATTAAAGC CCTGAACAGA CTAATAATGG NGITTCAAAA ATTGAATCAG TAATAAAA

SEO ID NO:500: (Length of Sequence = 474 Nucleotides)

TITIATITIT TICACIGITA CIGITITINA TCITIGATIG ATAAAAATGA AAATGCCAAA ATGAGGGITA GCTTAATITA AAGTATAAGC GTAGITAGCA GCTTTINCTA ATCACTCCTG TCCATTTAAA AAATAATCCT CATAGGAGTA TAAACAGAGG AAGGAGAAAT GGAGGATGGG CITAAGGAGA AGAGTATTTC ACAAATGTCT GCATAGCAAA TICAATTCAT CTACCTAGTA GCTCCTTCCG TGTTAACCTA CAGGTGTTCT CCCCTCCAAA AAAAAGCATC TTTTAGGAAG AAACCACCTT AACACTACCT TTTAGAAGATT GAACTTCCAG GGATAGGTTG TTTGAGAGAA TCACCAAAAG CCATTTTTAA ATGAATTTTT AAATTACGGC TTTCTCATTC CTTATAATAG TGTAGCAGCC ACCTTCCCTC TACTATGGAA CTTTTAACCA ATAATCCAAG TCCT

SEO ID NO:501: (Length of Sequence = 378 Nucleotides)

GIGGIGGCGG GCGCCIGACC TCGIGATCCG CCCGCCTCAG CCTCCCAAAG TGIIGGGATT ACAGGCGIGA GCACCGCACC CGGCCCTTGT GIACATTITT ATAAGAGAAT TITTITAGCT AGGAGITCAG AATTITIAAA GIACCATTIG AATGATCITA ATTITINCTIT CATGACAACA CATTCCAAAA TGAATCATGC TTATGIACTA AGAGGGAAAA TGIATTTAAG NIAAGGGIGA GAGACTTAAG TTATAGGIGA CCTTAGAGAC CTAAGGTGAG AGACTTGACA CATGGAAGGA GTAACATTAG GGICTACCTC TACCTCAATT TAGTTAGCGA TTTACTACAA TITCCAGAGCT AACAAAAGTA AAAATAAA

SEO ID NO:502: (Length of Sequence = 448 Nucleotides)

TTTTGGAGAT GGAGTCTTGC TCTGTTGCCC AGGCTGGAGT TCAATGGCAC AAACTCGGCT CACTGCAACC TCCGCCTCCC
AGGTTCAAGC AATTTTCCTG CCTCAGCCTC CCGAGTAGCT GGAATTACAG GCACACGCCA CCATGCCCAG CTAATTTTTG
TATTTTAGTA GAGACGGGGG TTTCACCATG TTGGCCAGGC TGGTCTCAAA CTCCTGAACT CAGGTGATCC ACTCCCTCGG
CCTCCCCAAAG GGTTGGGATT GCAGGCGTGA GCACCACGNC CAGCCATGAT CCTTAAACTT GTTTTAAGAG GTATAATAAC
TGGAAATCAT GATGCTCTTT AAGGAATACC AATTGGATGT ATTATTGATG TATTTAATTC CATCCATATG NAGTAGAAAC
AGTTTTCATT AGCAGAAGGC AATTATATTA TAGCTACACA ATATAAAG

SEO ID NO:503: (Length of Sequence = 446 Nucleotides)

CTACAGTACC CATCTCCATT TTCAGAGAGC TCCGATGGAA ATTTCTATGA ACTAATTCTC CTGCACATAC TTTGGTACAA GTGGGCTACT GGAGCCACCT TCCTTCGTTC AATCAAACAG CATTTATTCA GCTTATTTAA TGAACACTAT CCAAGATACT TGGGGGACAG AAATGAAAAG ATGGGGAGAC CTGTCAAACA TATGGTACTA TGTCTATGCA AAATAACATT GGAATGTAGA TTCACAGTGG AAGGCAGGGC AGGCATGGAA GAATTCTGAG AATGAGTGTG ACAGCTCCTA CCTGTAACAG CTCTTCAAGC TCCTGCTGGAA AGCGGTCAGA AGTGAAATTCC GGGATTTACA GAGCAGGTAG AGGGCATGCG GCCCAGCCCT CAAGCA

SEO ID NO:504: (Length of Sequence = 248 Nucleotides)

TTGCTCTTCT TTTCTACCAT GGGAACGTCC TTCTCAGGGG ATTTINAGGT CTCGGTGTTT CTGTGTTTCT NAATAGGCAG
TTTCTCGCTG TCGGCTAAGG GCTTATCCAG GNCAATATCC AGAGCCCTGT AGGGGTCGTT GGGGTCTTTG TCATCCTCGT
CGCTGGGCAG AGCATTCTCA GGCATCTCCT CTGTNACGAT GTCCACCTGC TGGGCAAGGG CGATGTCCTC GTCGCTCTCC
GTGGGCAA

SEQ ID NO:505: (Length of Sequence = 367 Nucleotides)

GCIATGITGC CCAGGCIGIT CTCAAACCCI TGAGCTCAAG CAGTCCTCTC ACCIGICTCC CAAAGINCIG GGATTACAGG
CATGAGCGAC TGINCIGGGC TTACTAAATT TTAAAAGATT TGIGITGAAC CATCIGCIGA TCATGGAGCA GCAGAGAAAT
TTATTGACAG ATTITCTAGG GTCATCACTG ATGACAATCT GNIGCCAGAA CAAGCCIGIA ATGCIGATGA AACATCACTG
TTCTGGCATT ATTGCTCCAG AAAGATACTG ACTACAGCTG ATGCAAAGGC CCCTGIAGGC AGIAAGGATG CCAAGGACAG
AATAACTGTT CIGGAATGIG CTAATAATGC AGCAGGCATT CAATAAG

SEO ID NO:506: (Length of Sequence = 419 Nucleotides)

ACACCTEGTE ACTITACCIA TECCTATCAA AAGCCTEAGG AAACAACCAG GTCCCCAGAT GAAGAAGATT ATGACTATGA
GTCTTATGAG AAGACCACCC GGACCTCAGA TGTGGGTGGC TATTACTATG AGAAGATAGA GAGAACCACA AAATCTCCAA
GTGACAGTGG CTACTCCTAT GAGACCATTG GGAAAACTAC CAAGACCCCT GAAGATGGTG ACTATTCCTA TGAAATTATT
GAGAAGACCA CACGGACCCC TGAAGAGGGT GGGTACTCAT ATGACATAAG TGAAAAGACC ACCAGCCCCC CCGAAGTGAG
TGGTTACAGC TATGAAAAGA CTGAGAGGTC TAGAAGGCTT CTGGGATGAC ATCAGCAATG GCTATGGATG GACTCTAAGG
ATGGTTGGCC ACACAACTT

SEO ID NO:507: (Length of Sequence = 417 Nucleotides)

GAAAACTATT TTACTTAAAA AATATTCTAT TACTTCAATG TCATGTCTGT TGAACGAGGA ACTCAACATG CTTATTINCC TTTGGTTCCA AGAAAAACCC AAGTCTAACC AAATGTATGC CACAAGGAAC TGCCAACTGG GTTAAAGCTT GGTATTTTCC TGGTTATCAC CCTATTTCCT GGTGTAGGAC CTGGGGTTTA ATAGAGACAT TTACATAAAA AAGGTATTTG GTTAAAACAA GAAATATGCA TGCNCTTCCT TACCACCTTC CTGGGAAAGA ACTGCTTTTT TTNCTTTCTT TCTGTGAATC TTGTTCAAGA CATCCTGTAG TTTAGATATA TGGGCGCTT CTTTTTTTACC CTCAAGCTTT TAGGTGACAC TTATAAAGGT GAGCATATCA TTCTATAAAAA TGGAAGA

SEO ID NO:508: (Length of Sequence = 308 Mucleotides)

CIGITTAGAA AAAAAAGIGC AGCICACIGI CAGCACICAT TGAATITIGC ATAAACAIGC TITITGAGGC TGAAGCAAAT CIGACIGATI TICAATGIGA AAAIAAAATA TAAAANCIGI TITITACAGIT ATTIATTAAC AGAACIAACA TCAGAATIAT TIGAATCACC AGAATAATCA ATTCIGGAAA AATCAGATTC ATCAGATTAA TCITTGGCCA ACAACIGITC AAGAACAATG TIAACATCIG CAIGGCAAIG CIACATITICC TAGGATTIGA CALITTCAGC AATTGAGGAA TTACTATA

SEO ID NO:509: (Length of Sequence = 370 Nucleotides)

TTTTTGAGAC GGAGITTCAC TCTTGTTGCC CAGGCTGGAG TGCAATGGCA TGATCTCGGC TCACCGCAAC CTCCGCCTCC
CGGGTTCAAG CGATTCTCCT GCCTCAGCCT CCCAAGTAGC TGGGATTACA GGCACGCGCC ACCACGCCTG GCTGATTTTN
TATTTTTAGT AGACACGGGT TTTCACCATG TTGGTCAGGC TGGTCTCAAA CTCCCGACCT CAAGTAGTCT GCCTGCCTCA
ACCTCCCAAAA GTGCTGGGAT TACAGGCGTG AGCACTTGCG CCTGGCCGTG ACTGATTTTT TTTCATGTAG AATTGTCAAC
ACGAGAGACTC ACAAGTGGAG CACTITGAAA GACCGTCGGT TGTGTGCACG

SEO ID NO:510: (Length of Sequence = 446 Mucleotides)

TCTTTCCTCT TACTTTCCTT CCTTCCCTCC TTTCATATGA GAGACTCTAT ATGGAAAAGG AAGCTGAAGT GGCCTGCACA
CGATATAGAA AAGCCATATT ACTTTCCTAA GACTGGTAAT CCGGCAATAC CTAATGCAGC ACATGGCTAG AGACTCCACA
TTTGCCCCAAC TTCTCTGCTC ATCATTTGCC ACTGTTCTGT AAATTTCCCA GTCCCCTCAC AGAAAGCACA TGGCACCATT
TAAAAATGGCT GCTCACTCTC TAAGGGAGGT CTCACAGGCT GGTAGTGAGC CCTGTCCCAA TAGTGAAGTT CTCCACAAAT
GGGGAGACTT CTCCCAGGAG GAGGGGAGGC CTGGAGATGG GCATGCAGTG GGCAATGTCA GCTGCCCTCC AGGTTCTTGC
TTGCCCTTTT TCCGCCCTGG GTCAGTATAC AAGCTTTCGG GGGACA

SEO ID NO:511: (Length of Sequence = 354 Nucleotides)

SEO ID NO:512: (Length of Sequence = 374 Nucleotides)

CATGUATATT ACAAAAAAGI TCCIGIACCA AAGITCITAT TAGACITTAT TITTIGITTIT TITAATITTIA AAATITTITIT
TGITTITATT TITATTITIT AAATTINCIC TCCIGGIGGI GACIGICATG TGATIGICIC AGITICIGGA CCAAACAAAC
ACACTAATAA TTITAAATCT GAAACAGIGA TIGICCCTIT NGGCTCATGI ATGIACAGGG TGATCAGAAG TGGTACCTGI
TAGCAAAAAGI GTCACGATGC TGCACCTCTA CCGAAACIGA TACCCACGAA CTACGGAATC TAAACAGACT ACACCCTGTA
ACTGCGTATT ACTGTCCACA ATGGGGATCT CCAAAGACAA AAGAGGTATG GAAA

SEO ID NO:513: (Length of Sequence = 463 Nucleotides)

ATCAGCAGAT TITNCTCIGG TGAATGICTA ATCAGTGTGA TITCCATAGG CTATACTTAC CTTTTGGGGG CTACTTGCCA
ATNATGITTG GTCAGTATCC TTGCAAACAA CAGAGTGACA GATTCTAAAA ATGACTTTGC AGGCCAGTAC TAAGAAAGAC
ACCAAGGTTC ATGGGCTTGC AAATAAAAAG TCCATAACTT CCCTGCCCTA CTTCACCAAG TGAAATCGAG TTCCTCACAC
TTCTGCACAC AGCTCTTCA GGATCTTCCC TTCCCTTCAA GGCTGTCTGA TGTTCAGTTT AATTTGATTG TATTTGTATA
AAGTGCTGAG TGTTGAGTCC TCAAAGAAAT TTACTTTCAG TCTAANGCCC CCTTGGGACA AGAAAGTGGC AACCAGGCAA
ATGATTGATT ACTTATTTGT TTGAGTATCA CTTTGTGATT GTCCCAGGGC TGTATTACAC ATA

SEO ID NO:514: (Length of Sequence = 396 Nucleotides)

CCAACCCAGA AAACGITTCC TEGETETETA CTAACAGTAA AATGTECTEA GCCCAAATTT TCTECTCTAA CATGGGTCCC ACGGACCTAT CAGTCTGCTC TEGEGTECTG ACCTECTGGG TCCTGAGCAG GGTCTTTCCC TAAGCATCAC TGTGGGTTTG GAGACAGCTG TAATGTGTGC AGCTGTCAGC AGAAAGTACA ATGCCACTGG GCTACATATG TCCATATCAT CCACCACCAT

TTCCCACTGT AAAACCAAAG GCTGCAACTG TGAACAAATG TGGACTTCCT CAAAGGACAA ATGAGGAGAC TGAAGGCTAC
ATTTCCTCCT TTGAGAACCC ATTAGAGAGT GTCTACAGTT ATACAACAGG TTTCTGCAAG ACCCTGTGGG TAACTT

- SEO ID NO:515: (Length of Sequence = 416 Nucleotides)
- ACAAAACAAA AAGTAGTAGC ATCTCTGTGA GAGGTACACA GITAGAAAAA TGATTCCACA CACGAGTAAA GAGATTTACC
 AGGAAGAGTC TTGTTTTCTA AAAGTTGATA CAACTAGTAG AAAAATACTT GTCAGTGGTA AATAGAGCAG AAGTAGAAAA
 AGCAGTTAAT CTATTAGATC AGATCAGAGT GTAAGGCAGG TATATCAGGC CAAAGGTGAT AAGACAGAGC AGAAATAAAG
 "TATTGTTAAT TCATGCATTT NCTGACTCAT TTATTTATAC ATTGATACTG TCACTTATAA ATCAAATCTT ACAGGTCAGG
 TTCTGTGCTA AGCTCAGGGG NTATAAAANG AAATANGTCA CTGCACTCGC CCTCACGGGG GCCCACCAGT ATAACTGGGT
 AGATAGTTCT ATAAAG

SEO ID NO:516: (Length of Sequence = 368 Nucleotides)

CCCATGGAGC TOGAGAACAT OGTAGOGAAC AOGGIGCTAC TOGAGGCCOG GGAAGGTGGC GGTGGAAATC GCAAAGGCAA
AAGCAAGAAA TGGCGGCAGA TGCTOCAGTT CCCTCACATC AGCCAGTGCG AAGAGTGCG GCTCAGCCTC GAGCGTGACT
ATCACAGCCT GTGCGAGCGG CANOCATTGG GCGCCTGCTG TTCCGAGAGT TCINIGCCAC GAGGCCGGAG CINAGCCGCT
GCGTCGCCTT CCTGGATGGG GTGGCCGAGT ATGAAGTGAC CCCGGATNAC AAGCCGAAGG CATGTGGGCC GCANTAACCG
CAGAATTTTC TNAGNCACAN GGGTCCTGAC CTCATCCCTG ACGTTCCC

SEQ ID NO:517: (Length of Sequence = 393 Nucleotides)

CCCAGCICCT GGAGAGCCAG CCCTGCAGGG TGGGCTGGGC GAGCCAAACT GCGTTCCTGG TGCAGGGCTT CGGGTCTCCC
TAACAGACCT TATACGCTGA CCGGCGGCG CCATGGCAGT GTCTCTTTGC TCAGACATCC AGGGACGACC ACATTCGTCC
AACAGCGGTC GCTCCACCAA TCCTGGGAGA AGGGAATCGT TTTCTCCGCG TGCCCTGTCA GCCGCTCATG GTGCCCAGAG
AAGTCAGGAA GAGAACCACC ATCAAGGTCC CAGGCTCTTT TTTTGTGACA AGGACTTAGA GGGGTTTGGG TCT

SEO ID NO:518: (Length of Sequence = 465 Nucleotides)

CCTTCTCTGC AGATAGAAGA GCCAGAATGG GAAAAGCGAA GATCCATCAA CCTGTCTGAG CTCATTGATG TTTACAGTGA
TGGTGTTGAA CTACTCCAGA TGGTGAAGGC ACCAGATTCC AACTGCAGCA ACCTTCTGAT TACAACCAGA CAAAGCCTTG
TNCTGCTTCG GGGGCAAAAT CTGACACCTT ACTGGGCATT GAGACTTCAA GGCCTGCGCA GCAGCCTACT CCTGGATATT
TCACTGATGA TCAGACATTA GACTTCCTTC TGCAGATACA GGATGGAGTT GGGATGAAAA AGATGATGGT TGTGGATGGT
GACTCTGGGC TCCATTGTTT GGAGTTACCG TGCTCCGTTG TCACATGAAA GAAAACGGCC AGCCACCTCA GCAGTTACTT
TCAGACCAGA AGTCTGTCTT TCCTCTTCTG GGGCCGAAGG CTTGTCAGGT TGCATCTTCC CAATT

SEQ ID NO:519: (Length of Sequence = 382 Nucleotides)

GGCCGICGGI AACAGAAAC TCAGIGCATA CITIGCIGIT GITAGGITGI CAATATAGIC TITCIGIAGG ATGGATAGCA
TGITTGAGAG GIGCCAAACA AGAACITIIG GGGITAGIAG TGIGICITGI GGAGGGIATI ACAGGACIGI GIAATTATAG
GACTCIAACT TGACATGGCI TGGCACCCAC TIGCAGCIAG TGGGIACAGG GIACAAAAGA TGITAGAGAA AAGCICIACA
GATTACGIAC TICTGIGICT TCGIATGCIC AACACTGICC TTTTGTCCTC CATGAAAGAT GAAGGAAGCA AATTTATGTA
TGINCITTCI TIGACCITCT TTAATCCTCT GATACTTTIT AGATTGCATG ATTTTACTAG GC

SEO ID NO:520: (Length of Sequence = 304 Nucleotides)

CCAAGACTGC TGATCTCTAA ACAAGCATCA AAACCCGAAG CTCATTAACA TCAGAGTGAG CTTCAATAAG GTGANCACTA
CAATGATGTA CAATTACATC CTAATANTTC ANTGCCCAAG AGCCCTGTAG AACTATTGCA AGGCCCAGGN TTATCACAGT
ATGCAAATGC ACTAGGAAAA TCATTACCTA TTTAGTCCCC TTTATTTTGG TGGGTTTAAC ATGAGAAGAG TAATCCATGC
TACAAGACGA GATTTCATTT TACAGCTGTA GTAGCCCAAGT GCATAAAAGC TTGANTCTGT CCCA

SEO ID NO:521: (Length of Sequence = 360 Nucleotides)

TTGAGACGGA GCTTTCCCTG TCACCCATGC TGGAGTGCAG TGGCGCTATC TCAGCTCACT ACAACCTCCA CCTCCCAGGT
CCAAGTGATT CTCCCGCCTC AGCCTCCCAA GCAGCTGGGA TTACAGGCGT GAGTCACCTG CCTCAGCCTC CCACAGTGCT
GGGATTACAG GTGTGAGCCA CTGCGCCAGG CCTCCCCAAGG TGTTGGGATT ACAGGCGTGA GCACCGCTCC GGGCCTCCCA
CAGTGCTAGG ATTACAGGTG TCAGCTGCTG CACCTGGCAA TTTTTTGATA TTAGGTCCCC TGAAGTCCAA AAAGAGATAT
ATGGCTTATT TGGTATAATG AAATCATACA GGAAGGCATT

SEQ ID NO:522: (Length of Sequence = 287 Nucleotides)

TTGAGGAAGT TCTGTTGCTG GTGAGGAAAT TCTNTTGAGT TCTGTAGGAA TTTTTATAGC TTGTTTTGCA TTCAGTTCTA
TCAACAAGCC AGCAGCAACT CAAAGGGAAG CCTCCTNCTG GCATATCAAT CACACAGGCA CATAGGATCA TATAGCATAT
AGGATCAGTC CCCAAGAAGAA CTATNGGGTN GGGGAGAGGT TTTTCTTCCA CTTCTTGGGN TTCAGTGACT TTGAGATGGA
CCTCTTTTTT CCNNTGGACA AAATGTCATC ACACCAACAT CTTATTG

<u>SEO ID NO:523:</u> (Length of Sequence = 318 Nucleotides)

CCTTGTCTCT ACTAAAAATA CAAAAATTAG CCGGGCATGG TGTCACGTGT CTGTNATCCC AGCTACTCGG GAGGCTGAGG CAGAAAAAATT GCTTGAACCT GGGGAGGCAGA GGTTGCAGAC AGCTGAGATC ACTCCATGC ACTCCAGCCT GGGCAACAAG AGCAAAAACTT TGTCTACAAG TCCTCCTACG CTGACAGGTC CTCACTCACC TGAATCTTTT ACGCCAGCAG CGTCTCTTCA CTGACGTNCT TCTNCATGCC GGAAATAGGA CCTTCCCTTG CCANCGGGCA GTGCTGGCTG CATGCAGTCG TTACTTTT

SEO ID NO:524: (Length of Sequence = 238 Nucleotides)

ATCTCATTGG AGCCAGGGIT CCAGTTCTCA TGCAAGTCGG CCACAGGAGC CACGGAACCG CAGTAGGATT TCTACTGTTA
TACAGCCCTT GAGGCAGAAT GCAGCAGAAG TTGTGGACCT TACCGTTGAT GAAGATGGTA AATTGAAGTA GTAACAGTAG
AAAATTATGA AAGGAGITTG ATAAAAGGAA ATCTCTTAAT ATGCTAGAAA CTCCTCCTGC TTACTGGTAA TATATTAT

SEO ID NO:525: (Length of Sequence = 168 Nucleotides)

CCAATGAGIG TGGACCCTAA ATTTAAACAG CTAAAGCTAT AGTCTAAGGA CAGTCTCAAA TAAATACCTT TGAATTGTCA
TATGGIGCCC AGGAGGGICT TGIGGAAAGG GTTTCATGGT AGTGAAAGAT GTAATANCTC TTTTTTCCTT TTAACCCTAA
GCCTGICC

SEO ID NO:526: (Length of Sequence = 387 Mucleotides)

GGAGGTCACA CGGTGAAACA GACACAGTTA TATACAACAG GGCAGGTTT TAAAAAGAGT TGCTCTCAGA CGCATTTTTC
CTGCTCCCTA AAAAGCCGAG GAAGATACTG GNTCCACAGA AAGAAAAAGGC AATGCCGTAA CATGAGGCCC TCATGGCCGC
ACCGTCCAGG GGAAGGGCTG TTAAAAACAC AAGTATTCTT GTGAAATACT TCGATCTGAG CATTAAGGCA GGTCTGCAGG
AGATCCGTCC TGGGGACTCG GACAGCAACG CTACCGGCTC CGAGAGGACA GTTAAATGTC GCCTCCCGGC AAGAGGGGCG
GAGAGATCAG ACAAGGAGTT GTTCCTGAGT TNAAACCTGC TACAACAGCA AACTCCAATA AACTCAA

SEQ ID NO:527: (Length of Sequence = 336 Nucleotides)

, ;i,

TTTGCAGITT TACATTCCCC TAGIACATCC CIGCITACIC GGGAGCACAA AGCTTGGITG TAAGAAATTG TGATTTGGAA
GTAGAGAAAA GCAAGGAAGT CCAACCTCAG GAGIGICTCT GITACIAAGA GGAGAGTGAG ATCCAGGGIG TGGGAGATGA
TCTGAAGGIC TATGGGTGGG GAGIGCCACA GGAAGAAGGG TTCTGGTCGG AGITAAAGGA GGATATATCT ATAINCTGGG
AGATGAGCTG AATTCAGAAC ACATGGAATG GGAACAATTC TCCCCATACT GCGTTTAAGC CAAATTAGGC TGGCATCCCC
CACCACGGCC AACTAA

SEO ID NO:528: (Length of Sequence = 482 Nucleotides)

TITTACTCTA GOGIGAGGAG GGGCCTCCT AAGGAAAGTC ATGCTGGGTA AACTGTGCGA TGTTACAGAG CACATTGAGT
CTGTGGTCAT CGIGGTTCTT CTATCTTCAC TGTCACCTGT ATCCTGTTAC ACATACTCAG TTCCTAATTG TAAGCTCAAT
TTTGGTATTA GCAAAAGCAT CTGTCAGTTT TTCCTCAATT ACTCACACCT CTTCTTGCCT AAATAAAACA AAGAAACAAA
GAAAACAAGT GTGGTGTCAT TACACGTCTC GGGAGTTCCT CGTCACTGAC TTTATATATA TANAANAAAG AATGCACATG
CGGGCCACGT TCACAGATAG ACAGATTCAC CCGAAATTGA GGAATGAGGG GCCTTAAAGG CTGCCGANAA NCAAAATGGG
GTGGAAATTA GCAANCGTTG TTTTCCGGTC AATTNCCAAT TGTGCACTGG CTGCGTTGAG ACAAGNCCAT CTTCCAATTT
CC

SEO ID NO:529: (Length of Sequence = 412 Nucleotides)

CTCTCAGACA GTATCCTCCT CGAAGCAGGA ATCCTAGTAA ATCTCATCTG CGGCATGCGA TTCCTAGTGC AGAGAGGGGA
CCTGGGTTAT TAGAAAGTCC TTCAATATTT AACTTCACTG CAGATCGATT AATTAATGGT GTCCGGAGTC CACAAACAAG
GCAAGCAGGT CAAACTAGAA CACGGATTCA AAACCCTTCA GCATATGCCA AGAGAGAGGC TGGGCCTGGG CGTGTGGAGC
CAGGCAGTCT CGAATCCTCT CCTGGTTTAG GGAGGGGAAG GAAGAATTCC TTTGGCTACC GGAAGAAAAG GGAGGAGAAG
TTTACAAGCA GCCAGACACA GTCTTNCAAC GNCACCAAAG CCTCCGTCGC CAAGCTTTCG AGCTGGGGGC TTTTCCAGCT
TTCCCTCCAT TA

SEO ID NO:530: (Length of Sequence = 301 Nucleotides)

ACTITITAAT AATAGICATI TAAAGIGGGI GAGATAATAT CICATIGIGG TITINATITG CATITCICIG AIGCITAGIG GIGITGAGCA TITIGINCATA TAACINCIGG CCATITGIAT GICITITITI TITITITITIT TITITITITIGA GAIGGAGICT CACITIGICA CCCAGGCIGG AGIGCAGIGG CGCAATCITG GCTTACIGCA ACCICCACIT TCIGGGITCA AGIGATICIC CIGCCICAGC CICCCAAGIA GCTGGGATTA CAGGNGCCCA CCACCACGCC CAGCTAATIT T

SEQ ID NO:531: (Length of Sequence = 312 Nucleotides)

CAGATGAGAC CAGGCCTTGA CAGTGGGGC AAGTCCTACC AACCTGCACA GCACATCCAG CAGGNCAACT GTGGCTCAGC
AGGTGCCAAA TGGAGCCCAT GGGCAGAAGA TGCCCACAGC GTTCCAGATG TGTGTGGTCT GAGAGATAAA AGGACACAGA
ACAAGATGAC TGTGCAAATA GCCAAGTGGT GGCAGAAGTT CTGCATTTCC AAGAGATGAT CCACTCAATA ATTTGACGAT
ACTAGTTGGC CAACATGCTC AGAGAAAACA GNCTTATCCA CATCTGGAGC CTCATTCTCT CTCAGGATCA TT

SEO ID NO:532: (Length of Sequence = 313 Nucleotides)

GCACAACTCT CGACCTTTGG GAGCAGCCAG GGAGGAGTCA CTGTCCCAGC CCCTGGCCT AGGCACAAAG GGGTGGGAGA
GACAGCTGGG CCAATATGGT CTATTACCGC CTGAAACCCC GCCGAACCAC CCTTAACTCT GCCTTCAGGC ATATCCCCCC
ACGTCCATGT CCAGGAGCCC CCCTACTGTC CTGGTCATCT GTGGCCCGGG GAATAATGGA GGAGATGGTC TGGTCTGTGC
TCGACACCTC AAACTCTTTG TGAGTATGTG GGGAGGGGGT GTGGGGGAGG AGGCCGTNYG GGCTCTGGGA TCT

SEO IP NO:533: (Length of Sequence = 376 Nucleotides)

GTAATTCCAT GTGGCTGACT GGGTAACAGA TTTGAAGGGT ATCACAGACC TTCATGTTGT AGCTCATCGC AGTGTATTGT
TTGTTGCTTG TCTCTGTCTC CCGTTGTATT GCCATCCTCA AGGGCAAAGA CTGCATCTTT GTATTCCCAG CTCCTAGGCC
TGAGTCAGGC ACATAGTAGG AATTCAGAAA GTATGTTTTG GATGTAACAT TCCTCCTTTT TCCTGGACAA AATGGCCTTT
TGTCCGGTGC ATTGTCCTTT CCATAGAGGA GGGGTTGGGG CAGGATTGIN AGATGACTGT GTTTGAATCT TCAGTTAGCT
AAGACAAGGA TACGINTTTT CCATGGTGCA AATCTAAAGG GTTCTAGTGA GGTGGTTC

SEO ID NO:534: (Length of Sequence = 374 Nucleotides)

TITITITITI GICCAAGGIT TATCAAATTA ATTGATTITG GGGGGCAAGA TAAAAATTIT NATITGATTA ACTITCICIA
TIGGITITIG TITICAATTI CATTTATITC TICTITTATC TITIATAATGI NCITACATCI GCITGGITIG GGCIGGCAC
AGGGGCTCAT GCCTGIAATC CCAGTACTIT GGGAGGCCAA GGTGGGCAGA TCACTTGAGA CCAGGAGTIT GAGACCAGCC
TGGCCAACAT GGGGAAACCC CGTCTCTGCT AGAAATATAG AAATTGGCCA GGTGGGTGG CCAGCACCTG TGATCCTAGC
TACTCGAGAG GCTGAGGCAG GAGAATGGCG TGAACCCGG AGGCAGAGC TTGC

SEO ID NO:535: (Length of Sequence = 433 Nucleotides)

TGCCGACTGA TTCCAAGTCC CCAGGAGGC TGTGAATGCT AATAGATATT TGGGGTTTAT CTACATGGAT AAATCAGAAT
TGTTAACATT ATTTATAAAG ATAATACITA CATAATTTIN AAATTCACAA AGATTGITTG GCTTAATGAT TTCTAAATGT
ATGCAATATA ACATTAGGCG GCTTTTATTA ATTCTATTTA TGTAATGGAA AAGCTCAATT CAGCAAAAAA CAGATCTGAT
GGGATTTGGT TATTCTCTAC CTGATCAGAA CAAAGCCTTA CTTTACATTC CTGACTACCG ATTGGCTGAG GGATTGTCTA
ATAGAATGGA GCTTTCTTTT GAGCGGTATC CATGTGTACA AAATTGGGCT GCTTTACCTG TGACCCACGG ATTGCTGGAG
GAGCTTG-AA ATGTAGTCAG CCGTTTCTTT TGG

SEO ID NO:536: (Length of Sequence = 438 Nucleotides)

GATGAATTAA GAGGGAAATT TATAAAGTAA AATCTTTAGC GCTGTTGATC AAAGAGTTCC AGGCCGGGG TGGTGGCTCA
TGCCTGTAAT CCCAGCACTT TGGGAGGCTG AGGTGGGCAG ATCACGAGGT CAGGAGATCA AGACCATCCT AACACGGTGA
AACCCCATCT CTACTAAAAA TACAAAAAAT TAGCCGGGCA TGGTGGCAGG TGCCTGTAGT CCCAGCTATT TGGGAGGCTG
AGGCAGAAGA GGAATTCCTG CAGCCCGGG GATCCACTAG TTCTAGAGCG GCCGCCACCG CGGTTGGAGC TCCAGCTTTT
TTGTTCCCTT TAGTGAGGGT TAATTTCGAG CTTGGCGTAA ATCATGGGTC ATAGCTGTTT TCCTGTGTGA AATTGTTATC
CGNTCACAAT TCCACACAAC ATACGAGCCG GAAAGCAT

SEO ID NO:537: (Length of Sequence = 316 Mucleotides)

TAGTAGCACT AAAGCCCCGT TTTGGTCACA CTCTCACCTA GGTGAGAACC TGACCAAAAA TGTGGAATTA TTAAACAAAA
TGATGGGAAG CCAATGINCT GAAACTGAGC TCTTGCACTA GGCCCCCACA GACCAAATTA AAATGGAGTC ACTAGTGCTA
AATGCTTTGG AGTCAAACAG AAATGTTAAA GAAGATAGAT CCCAAAAACAG AGCAGTGTTT TATTTTTTCTC CAGAAAACAG
GAGATTCCAG CATAATAAGA AAGTCTCCTC TGTTGTAACC CTTACAAAAA AGTAACCTGA AGTAACCATT TTTTTT

SEO ID NO:538: (Length of Sequence = 303 Nucleotides)

ATCITCATGG GCGTCCTAAC TGIAACAAAA ACCCCACAAT TTGAACAGAA GAACAGAAGT ATCTGGTTAC AGAAGTGCAT TCATACATTT CACAAATGTT TCAGIATCCT CTTCTCCCCG ACCCCAGCAT GAGCTTTAAT TGGATGTATT TATTCTTTCA CCAGCATGCC CATGAAGGNG CTAAGGAAAA CATTTACCAA GTCTGTTTCA AAATCTGTCC TTGGCATATC AAACTTTTTC TCTTCCTTTT TCATGCTTTT TTTTAAAAAA AAAAACAGGA GAAAGCGAAT AGAGAGGAAA GAG

SEO ID NO:539: (Length of Sequence - 162 Nucleotides)

, i. ,

CATGICATAG TGGCCIGCIC TCCIAACACA GCACAATTIA GGGCATATIT TCATGATGGT CIATCACTGG ATTACAACAC
ATCICITCAT TAAAGICITG GGAAAGAGGC TICAACITIN CIGIGITGAG AAAACITCAC AGGIGIGTAA AGITTGATCA
GIATGIATAA TATATTINAT TACATATATI TNATTITNAT TITTCATTIT TTIGCATACA TAGCAGGIGT ATATACITAT
GGGTTATATG AGATATTITG ATAAAGGCAT GCAATGIGTA ATAATCACAT CAGGGTAAAT GCAGTATCTA TCCATCACCC
CAAGCATTIA TCCTTTGIGT TACATACAGT CCAATTACAC TC

- SEO ID NO:540: (Length of Sequence = 416 Nucleotides)
- " CACCAGGGAG AACCAATACA ACAGAAAAAA AGCAGAGAAC AGCTATGTGT CCTGCCAGGT CTACCAAAGA TAGTCATCCA
 AACATGAACA GATGAGAAGG CTGTTTTTCA AGAAGGTGAA AGTGACAGAN TATTCAATGA ATCTGAACAC ATGAAGATAC
 TGAGACACCA GTAGTTCAGC AATAAGTGGA GAGAAAACTA AGCAAATGAG AAACTTAGGA ACAATTATGC AGCAAAGAAC
 AACTGGATAA GCTGAAAAGT GTTTAAAGAT GCTGCCGTAA ACACTAAGTA TCACAATCAA ATTCTGATTT GTAAAAATAG
 AGGTATGGGA AGGGTACANG TATGTTTGTG GGGCAAAATG GTGAGGAGGG CTTAAACCCT CTTCTTCCTT AATGAGGAAT
 TAAATAATCC CATTAA

SEQ ID NO:541: (Length of Sequence = 341 Mucleotides)

GAAATACTIC CAGGCCITCG AAAGGCCATC CITIGGACAC ATGIAAAAAG CIGICTIGIT GGCCCGITAT TCCCACTGAC

CCGICTGAGI GATCACCCAG GAGCGCGGCG GCAGCAAGCA GAGCTCACCG GATTTGGGAC AAGGATTITA AAGGCAGCTA

CAAAGCTGAG CICIATTIGC TGATGATAGI CICIGITCAG CIGITTAAAA TGACTGICTG ACTCACCATG GIAATTTINC

ACAAATTAAA AACACATTIT GGGITGIGCA ACAGIGGITC TCATCITTCC AGGCAGGCAG ATTATTITAA TGCTGITTAT

ACAGGGAATT GGGACTCTCG G

SEQ ID NO:542: (Length of Sequence = 334 Nucleotides)

TTGTTGTTC CTACCTTAAC CAATACCTCC TGGAAAAAAG AGGTATTGGT ATAAAAATAA ACCATACCCA AACATTCCCA CAACATGACC TTAATAAGCT GGTGCACAGT AGATTATGGC AGAGGAAAGA AAATTGACTT TAGAATTAGA GAAACTTAGG TTCAAATCTC AGCTCTGTCA TGCTTTGGTT GACCTTCAGT AAGTCCCATT TNCTTCATCT GTAAAATCGG AATAACATCT ACTCCACAGC ATCATTAGAA AGATTAAATA GTGGCTGGGC ATGGTGGCTC ATGNCTGTAA TCCCAGCACT TTTTGGGGGAGG CTGAAGGTGGG GCGG

SEO ID NO:543: (Length of Sequence = 350 Nucleotides)

ATTIGITIGC AATIGACAAC ACCICATIAA TIGIAAGCCC AGIGACACIG CITGCIGITI CAAGICACIT TIAAATIACA CACGIGCIAC TIAAATCTIAA AAGCAAAATT AAACATIGGA CIGGITIACA TITCAAGCTA CAATATGGAA CCATTGIATT TGGAGGAATG AGITTAATAT GCATTGIAAA ATAAAATTAG GGGGIACTTI GCATTCACAG CGGCTTATGI AATIAGGITC AGICAACTGI AATGITICAG GTTAATGICT TCCATGGATG TATGCTGIGI AAATAGTGAA CITACATATC CCTTAATACA TCTGAATTAT TACATAAATC CTTAATATTA

- SEO ID NO:544: (Length of Sequence = 328 Nucleotides)
- GEGAGACGAG AACTCTTGAG ATCCGGGGTC ACCTGINAGT CGCTGGACCC AAGGGGGAAG CGTCTTGATT CCTGGAGGAA
 ATCTCCGAAG TGATGTGIAA CCCTGTGTGT CGCCTGCACT TCGGCCGCAA CTGCCTTTGG TTCAGTCCCC TGTTCCTGTA
 GGAGGCGGGG ATCATGIAAC AGTGGAGCAC ATCGCTCCCG GCTTGGACGC CTTINACCTT TAAGTGTTCC TGATTTAGTT
 TGGCTTTGGG TCTACCAAGA ATTCTAGTCA GTTAACTAGC TTTTTTAAGCC AGGTTCCTGA ATTTGGTAGG CATGGACACT
 CCCCAGTAG

SEO ID NO:545: (Length of Sequence = 342 Nucleotides)

GGGCTATTAC CICIGGGCAC TGGGAAAACT GGGAGACGG ACAAGGGGT ACCAATITIT CAGIGTATGC CCTTTTCGAA
GTGTTAAACT TTTTTTTTT TTTTTTGAGA CAGGNTCTCA CTCTGTTGCC CTGCTGGAGT GCAATGGTGA GATCGTAACT
CACTAAAGCC TCAACTTCCT CGGCTCAAGC AATCCTCTCA CCTCAGCCTC CTGAGAAGCT GGGACTACAA GTNTGTGCCA
CCATACCTGG NIAATINITA AAGITTTTGT AAAGATGGGG GTTTTCCGAT GTTGCCCAAG CTAGTCTCAA ACTNCTGGGC
TCAAGTGATT TGCCCACCTT GG

SEO ID NO:546: (Length of Sequence = 280 Nucleotides)

CTOGTAATGC CAGCATTITG GGAGGCTTGA GGCGGGAGGN TCCCTTGAGC CGAGGAGTTC GAGATCAGCC TATGCAACAC AGTGAAGACCC CTATNICTAT TINATITAAA AAAAAAAAA AAAGGGGGTC ACGTTTACTG CCACCATCCC AGGCAGAAAG ATGAAGCCTA GAGCCTCTCA CTGCTTCCTA GTGGGTCTTG GGTGTAATT TGCTGTCTTG GGTATATTTT TTGGCAGAAA GCATCTGGCA TCAGGCACTG GTTCTCAAAG TCGGGCCCCC

SEQ ID NO:547: (Length of Sequence = 298 Nucleotides)

CTAAAGAGTT TCACATAGTG GCTCAGTCCA GCCTTGTGGG GATCTTGCCG GGGCCTGGGG CCGGTGGTCC GGGGCCTAGG GGGATGCCTN ACCAACAGAG GCTCTNCAGG CTCTGAAGAT AAGCTGAGGG CAACAGTGGA CAGAGGGGGC TGAACTTGCC TCAAGGAGGC TCTTATTCAA GAGCAAGTCT TGCTGGCTTC TNCTGAGGCT GGGGACCACG TGGCCCTTTG GCCAGCCAGG ACCAGCAGCN CTNACCACCT GCTGAGGGGC AGTTTGGGTC AGGGGGGNCA CATAGAGG

SEO ID NO:548: (Length of Sequence = 311 No: eotides)

GAGACAGGC TGITTCCTGC ACTACACTGG TCATCTGACC ACCTTTCTGC AATGCTAAGA AGGTATTCTT TGACCAAACA
GCAGTCCACA TACAAGTTTA AAAGGGGCCC TGTTTATGTA GGAACAACAC TGAGGTGGTG CGTAGCAGGT ACAAGACGCC
CAAATATTTC CAGTTTATCT TACGGCTGGA CTCCTATTCT CCCACACTGT TTCCTAAAGA AGGTCCACAT TATTTTGGNT
ACTAGCCTAG TTTAAGTGGA GATACTGTGG GCAACTTNAA AGAAATGACA TCAGGCACAC AGGCTGAGCT T

<u>SEO ID NO:549:</u> (Length of Sequence = 387 Nucleotides)

TITATITIGG TGTAAAGACA GGAAGCTGGA AAATACACTG TATITAAAAT TINCITGGIT CCCCCTCACA TIGTGGAAAC CCCCTCCCCC CAGAGCTAAT CTGTICAAAC TCAAATACTI AAAAATTACA GCAGCCAAAC AAAAGCATGG GGGAAAAAAA AACAAAAAACA AAAACCAGAT GGAGAAGGTA GCCTGGGCCCA GTAGTGTCAC TTGGTGTGGA CGACTGAGGT GCTGAACAGG AGCTTCTGIT TCTGTTTTT TCTTTTCTTT CCTCCTTTCT CTTCAGAGAG GGGATCTNGA AGTAGCTGGG TGTGTCCAGT TTCATGAAGG CTGCTTCAAT AGCTTGGCTG AAGGAATTTT GAAAACTNGG CACAGGAACA CGGGTTT

SEO ID NO:550: (Length of Sequence = 377 Nucleotides)

CACCCCCAAC TCTTCACCAA GTAGGGGCCC TGGCTTGCAA TTGCAGAAGA GCTTTCCCAT CCCTGGGTGA GCATACCTAC TGGTAGTGGC TCCGTGATTC CCTGGGGAAGA GACCAAGA GGTAACCAAC CAACCCTGTG CTACTGCTAT GACCACAGTT CTGCTTCTGC TGCCCTCAAA CTGGGGAAGA AACAAAGAGC CTGAGGGCTT TACTCACGCT TCTAGCACTA CGCAGTCACC ATATAAAGAG GAGCCCAGTC TCTCTTCCTT GTGAACCCTT GACCCCCAAC TCTTCACCAA GTGGGGCCCC CAGCTTGGGC CAGCAGCACA GTGGCCCCCAA CCCCTAGGCT GAACATTCCA GTAGCAGCTG CTCCGCG

SEO ID NO:551: (Length of Sequence = 320 Nucleotides)

GAGTTINIGG IGAGCCGAGA ICACGCCATI GCACTCCAGC CIGAGCAAIC AGAACGGICC GGCICCIGII GCIGAGGAAG CAGCTCIGGA IGACCIICAI GAIGAAAITI GCAGCCICGC GCICAGICAI GTIVFFFCIF AACCIGGGC IGGAGAAGAG GCTGTGTCAG GGCTGCCATG GGCAGGGCCG TGCTGGCTCC CTGGCCCCAGT GGGAGGAGGG TCTTCCATGG GGACGGACTT CAGCTGAGAG CCATGCCCTG GGAAATGTAC CTTTGGGGTC CACATGTTGG AAGATGGGGT GCTGTGAAGG CCACACC

SEO ID NO:552: (Length of Sequence = 334 Nucleotides)

ACAAACTGAC AAGAGAAAAC AAAGAATTCT TTGGTGATCT GGACACGCTG ATGGGGCCTC TGACCCAGCA CAGCAGCAIG
ACCAATCITG TCCGCTACGT TCGCCAAGGA CTGINTIGGC TGCGCATCGA TGCCCACTTG TTGTAGTGGG TGTTCTCAGA
TCTCTAGCAT CACGACCCAT CACTCTACCT CTACCAGCGC ACTGATGGTC ACTGGTGGAA CTCCACTCAC TGGGGAACGT
TCTCTTTGGT TATGTTTGTT TTTATGCTTC TTTTGTTATC TGTAAAAAAAC AGAAGTCATT GTAAGTTGAC ACTACAACTT
AAGGGCAGTG TACG

SEO ID NO:553: (Length of Sequence = 371 Nucleotides)

GAAAAGGGGA AAAATCACAA TATGTGTTCT AGACAATATT GGTTTAGATT TTTTAAAGAT CTAAAATCA ATTATGGAAA

GCCAGCAGCC TGATCCAGTT ACTGTGACTA AAGCAATTGT CAGACCATCT CTAGTCAACC CCTTATGGGT TTTGCAATGT

GTCTACCCCA ATTTTGGATC AGGAGGGGTC AAACAGAAAT ACAGCAATGT GATTAANCTG CTCCTTTGCA AACAATATGA

AAAGGTTTGT NCTTTCAAAG TAGATTCTAA CAAATCGTCT GCTCACTGTG GGGTAGCAAA GNGAGAAAAG CAAATCTTTC

TATTAGTCTC AAGCAAGTCT TCAGATTTAC ACACAATCTA ATGGAGGCAT C

SEO ID NO:554: (Length of Sequence = 331 Nucleotides)

TTATGACTIT TITCAATAAG GCTATTGTAT CAGCCTGTNC TCTCGCTGCT AATAACGACA TACCCAAGAC TGGGTAATTT

ATAAAGGAAA GAGGTTTTAT GGACTCACAG TTCCACATGG CTGGGGAGGT CTCACAATCA TGGCGGAGG CAAAGGAGAA

GCAGAGTCAC ATCTTACATG GCAGTAGGCA AGAAGAGCAT GTGCAGGGGA ACTCCCCTTT ATAAAACCAT CAGATCTAGT

GAGATTTATT CACTATCAAT GAGAGGCAGC ATGGGAAAAA CCTGCCCCTC ATGATTCAAT TACTTCCCAT TAGGTCCCIN

CCACAATACA T

SEO ID NO:555: (Length of Sequence = 305 Nucleotides)

GCTGGGACTA CAGGCGCCCG CCACCACGCC CGGCTAATIT TITGTATTIT TAGTAGAGAT GGGGTTTCAC CATGTTGGCC

AGGATGGTCT CGATTTCCTG ACCTCATGAT CTGCCCGCCT CGACCTCCCA AAGTGCTGGG ATTACAGGCG TGAGCACCGC

GCCCAGCCCA ACACATGGTA TITTCTGTCA TITTCATTTA GTCTTCTGGT TGCTGTGTGA TGGTCTCAGG CTTTATTTAC

ATTTCTCCGA TTACTAACAG ACTTGAACAT TTCAGCACAC TTTTTAGGTT ATTGAATAAC CCCTA

SEO ID NO:556: (Length of Sequence = 318 Nucleotides)

CITITITIGET GATINCTAAG CICIGITIIN CITATCCTAT ATATATATGT GGITGGITT NATTITAGGA TITTAAGGIT

ATCCCTAATA AATTITGAGA TGIGITCCAT AGCIAGCCIG TIGAGATCIT TINATATCAA AAGITAATAT CIGIGGATTI

MIAATCATTC TITCIACATA TITAACAAAG TCATTAGCAA AATATTGAAC AAAACCIGIT ATTCATATCC TIAGATACAG

AACATCAATA TCCTGAGATA CAGIACATCA TCAAAATGIG GICCCCAAAT GNGCAGCAAT TAGCATCATG TGGGAGCT

SEO ID NO:557: (Length of Sequence = 349 Nucleotides)

GRAAGCAATG TGCTTCTTCT TAACAGAGAT ACTGCACTAT TCTCTATGTA TACTCACTTG ATGGCATGGT ACATGTCCTC

CAGGATGTCT TGCTCAAAGT CCTTGCCTCC ATTCACACCT TTCAGATTTT TGCGAAACTC CTAGAGACAG GCCAGTAAGT

TTTTTCCCCCT TGTGTCAACA CTGAAGCCCC ACCTAAGGAA CTCTTGGGTT TTCAGTAAAT AGGACTTAGG AAAAGGTAAG

CGAAAAAAACC CACTTCCCCA CCCCAGTCCC TTTTCTAGGT TTGGGCCAGC CCTTCCTTGA TTCCCTTGGA CAGAACCCCA

TCCATCAGGC CCACTGGAAT CCTATGTCC

SEO ID NO:558: (Length of Sequence = 279 Nucleotides)

GGGCCAGGCG CTGTGGCTCA CGCCTGTAAT CCTAGCACTT TGGGAGGCCA AGGTGGGCAG ATCACCTGAG ATCAGGAGTT CAAGACCAGC CTGGCCATGT TGAAACCCCA TCTTTACTTG TAATACAAAA ATTAGCTGGG CGTGGTGGTG TGCGCCTATA ATCCCAGCTG CTTGGGAGGC TGAGACAGGA GAACCTCTTG AACCCGGGAG GCAGAGGTTN CAGTGAGCCA AGACTGCACC ACTGCATTCC AGCCTGGGGG ACAGAGTGAA ACTGTGTCT

SEO ID NO:559: (Length of Sequence = 278 Nucleotides)

GAGAAAGCCA AGAGCCATCT GGAGGTGCCG CTGGAGGAGA ACGTGAACCG CCGCNTGCTG GAGGAGGGCA GCGTGGAGGC GCGCACCATC GAGGACGCCA TTGCAGTGCT CAGCGTGGCG GAGGAGGCGG CCGACCGGCA CCCAGAAAGA CGCATGCGGG CAGCCTTCAC AGCCTTTNAG GAAGCCCAGC TGCCGCGCT CAAACAAGAG AACCCCAACA TGCGGCTNTC GCAGCTGAAA CAGCTCTTCA AGAAGGAGTG GCTCCGCTCT CCTGACAA

SEO ID NO:560: (Length of Sequence = 304 Nucleotides)

CANATGITAT TEGAAGITAT CIAGAAGGCT CAGTAACCAG AACTICCITT CATTCIGCT TICTITITCT TITTITTITT
CITCIGAGAC AGICIGGCIC TGICICCCAG GCIGGAGIGC AATGGIGIAA TCICAGCICA TIGCAACCIC TGCIGCCCGG
GITTGIGČAA TICICCIGCC TCAGCCICCC GAGTAGCGGG ATTACAGGCA CGIGCCACCA CACCIGGCIA ATTITTITTIT
TIGIATTITT AGTAGAGCCG GGGITTICAC CATGITGGCC AGGCTAGITT CAAA

<u>SPO ID NO:561:</u> (Length of Sequence = 323 Nucleotides)

GATGGTAAAC ATAAACCCAA ATATATCTGT AATTACATTA AGTGCAAGTG AACCAAAACA GITCAGATAA AAGACAGTAC CTATTTCATA GCATTATGAC TATCATGAGG TAATATATGT AGAGATTAGA GIACACATGT CATATTAGGA GGTGTGCAAT AAATGATACT TTATTCTGAA GATTAACATA ATTCATACTT AAAAGGATCA AGAACTAGAA TATTAAAAAA NTAGAATGTG AATGTTTCTG CAAGTTTTGA TAAGAACAAG CCCATAAATT AATCTCTAAT TTGCTACATT TAGGGAATAT GGGTAATGAC TAC

SEQ ID NO:562: (Length of Sequence = 214 Mucleotides)

TCTAATNAGG CCCTGCGTGC TGTGTCATCC CATGGCGGAA GAAGGAAGGG CAAGAGTGGG TGAGATTGTN AGCACGAGAG AAGGCTGAAC TTCATATTTT AACAACCCAC TTTCATGATT ATNATAATCT TCGCATTTAT TTTTTTCGGT CTCTTCATGT NCTCTAACTT TTCTCTGGGN TTTTGGTCTT TTGCTTCTTC ATTTTTAGAA GCTC

SEQ ID NO:563: (Length of Sequence = 358 Mucleotides)

TITITITITAT GAGAAACAGA AGCTGAATAT CCTGATTIGA TITGCCACAC AGGCGITCAA TGGCTAGCA GIGCTAAAGA
TITATTITTA TITTTITGGG CTCTGGGCTG ACATTGGAAA TITINCTGAA TGAGAAAAAC CATCCTCAAC CACTGITTIT
TAACACTGAG TAACTITGGA AATTAACTIT TGCCACAGAC TIGAAAATGI TICTTAATGA ATTTGACCTG AAATTACAAG
GTACAACAAC ATAATATGGI AAATTCATTI CAATAAAAAC TAAAACTTAA GATTGICAAG CTGCTTTATA TACTTNCTGT
GCTATGAGAA GTCAAAACAG CGCTGTATTG CCCAAATCC

SEQ ID NO:564: (Length of Sequence = 405 Nucleotides)

ATGRACIGIG TGTTTCATAC ACATGITTCC TTTAGTCTTA AAATCTGGCT CATGGGGTAA ACACTATTAT AATCTCCATC CTCCAGATGA GGAAAGTGAG ACTTAGAGGT TAAGTACATT TTAGGATAAA GTAGGGTATT TCGATAAATGT GTTTCTGGTC TCTGAGGACT ACACTCCCAG GCTGCTGGGG ATACAAAATA CCCT117C1TT ACCATAGGAG CACTTGGGTA GAATATTTGC AGAAACAATA AACTCGCTGA TATTTAAAGT TCTCTTCTGC TCTGACATTC TATAATTTGA TTGACCCTCT

TIGCATTIAA TIAIGITGAT TITCCITICT ACCCCITGCT TAGCIAAAAA TATACCCCIT CINIGICCAT GGACAGGAGG ATGGG

SEQ ID NO:565: (Length of Sequence = 196 Nucleotides)

CATCCACATC AGGCAAAGGC AAAGCAGGAC CTGAACTTCC CACCCCAAGC CCTACATCCA TGCAAGCCAG ACCAGACTGG GTCAGAGGCT AGAAGGGNGC TCACAGGNTT GCCTGGGGAA GCCTGGGCCC AAAACCTGGC CCTNGCTCCA GCCCAGAGNA CCCACCTGGG CATNAGACTT GCGCCAGCGT AGGGT

SEO ID NO:566: (Length of Sequence = 275 Nucleotides)

TIGGAAAAAA GAGAAAAAA AATICIGCIT CATITACGAA TGITGCCAAA GGAGGCAAGI TITCAACIGA AAACAAAACA
TAAAGGICIA TGIGGATGCA GCCAAATGIT TCICCATITA GAAAATCATC ATAAAAGGIG GCAGCACIIT TTITGCITGI
TAACIATATI ACITATAACI GGCIGCACCA ACATITCATC TCAATTITIG GAGIGITICI TCIGATCAAT CCTAAAAGCA
ACACAATCAT TITAGAGGIT GCAGACTACA ACAGC

SEQ ID NO:567: (Length of Sequence = 349 Nucleotides)

CECTOSINIG TOCCACACAA ATGITIAAGA AGICACIGCA ATGITACTOCC CGGCTCTGAT GAAAAGAAGC CCCTGGCACA
AAAGATTOCA GIGCOCCTGA AGAGGCTCCC TTCCTCCTGT GGGCTCTCCT AGAAAACCAG CGGGACGGCC TCCCTGCTGA
TACOGTCTAT AACCTTAGGG GGNCCTCGGG CAGGCAAACT CATCTCGGTG ATGCTGTAG ATGCTAACAC TGGCCAATTC
AATGCCACAC CTACTGGTTA CCCTTTGAGG GCATTTCTCC AGACAGAAGC CCCTTGAAGC CTAGGTAGGG CAGGATCAGA
GATACAACCC GTGTTTGTCT CGAAGGGCT

SEO ID NO:568: (Length of Sequence = 368 Nucleotides)

CIGGIAACIT CCCGATIGGN TITCCCCGCC TCANCCCITT CCCAGGGCIA TICICCTCCC ACCIGCIGCC AGGCCITTCC
CIGGCCATCC TGIGIIAAAT GICAICCCGC CCCIACIGIT AIGITCICCA CAGCACITGA ACACGACCCA ACATGCCITT
TCACITCAAG GITTATTCIT CIATTAGITT TCCCAGAGIC TGCTTCCCIA GIGICCATCI CCCCIGCTCG AATGCCICIT
GAGAGCCAGI GCTIGIATIT TGGTCCINGI GGIATGGGCC TGGCACATAG TAGGCAGICA GCAGATATIT ATGGAACAAA
CAAATGAATT TGTGTGACIA TAGITCATIG TTCATAGITC ATTCATAG

SEO ID NO:569: (Length of Sequence = 328 Nucleotides)

TETCACITAA TECACAGCIE GEGCTCAGGA CACAGCITIE CACACCCIAA GINCTCAATA AATECTAGCI CAGGGCAGAG CTTTGCATAC CCTAAGTACT CAATAAATGC TAGCTCAGGG CAGAGCTTIE CATACCCTAA GIACTCAATA AATECTAGCT CAGGGCAGAG CTTTGCATAC CCTAAGTGCT CAATAAATGC TAGCTCAGGG CAGAGCTTIE CATACCCTAA GIGCTCAATA AATECTAGCT CAGGGCAGAG CTTTGCATAC CCTAAGTACT CAATAAATGC TAGCTCAGAG ACAGAGCTTI GCATACCCTA AGGIGCTC

SEO ID NO:570: (Length of Sequence = 313 Nucleotides)

CCCTAAAAGG CAGAGTGTCT TCTTACCTCC ACACAACCAC GCTAGCTCTA TAGCAGTGGT TCTTAACCAG ATTGAAATGG CTGAAATGAC AGACATATAT TTCAGAACCT GGATGGGAAG AAAGCTCAAT GAGATAGAGG AGAAGGTTGA AACGCATCCA AGTAAAGCAG TAAAATGATC CAAGAGTTGA AAGATGACTT AGCCATTTTA AGAAAGAACC AAACAGAACT TCTGGAAATA AAAAAAAATC ACTACAGGAA TTTCATAATG CAATTGGAAG CAATTGGAACC AAACAGAACC AATCTGAAGGC AAA

SEO ID NO:571: (Length of Sequence = 338 Nucleotides)

AGGAAACAG GGGTCTCAAT TCTGTACGAA AGAGGAGGGT GTTTTACTTC CTGGAATTAT AGAGGCCAGA GGTGTCTCTT
TTTCAATTTA TTGGGAAGGT TTATTTTAAT ATGGACTTAG AAATAAATAA CTTATTAAAG TGAAGGTTCA CCTGGAGCCT
TAGGCTGGCT GCTAAGTGTG AGTCTGGGCT GTTGAAGGGA CTGTNCTGTT CTNCTGGGTC TCTGTAGGAG TTTGAAGGAG
AAGACTGGCC CCAAAGGGTG TTTGAACAGG TTAGATGTGC CCATTGGTTA GAACTTACTT GGATAGGGAG AAGGCNTCTA
GGGCGTATCC ACAAACTT

SEO ID NO:572: (Length of Sequence = 375 Nucleotides)

CTATTTCCAG AAGTGACAGC ACAAGTCTGA GTTGCTGTTT GGTCTGGTGA CCTCAGACAC ACTAATTTGA ATTGAAAGCT
AAGAGTAAAA ATTINCTGGT TACAGGCGAG TCATACTCTT GCAAGTAGTT AGCAAAGGGA GGCCCAAATT CTCAAGGTTG
TTGATGGGGA ACTTGCCACT AAGAGAAGGC AGAGAGGTCC CTAGTGGGTA TATTTNCTGC CAAGCCACTT GCCAAAGAAG
AGGAACCACA GAAAGAGAGA CATCATGACC NGGAGAAAAA TGTGACTAGA CATGCTAACC TCCAGGTNIT TATATATGAC
TTGAGTCTGC TGTAATTGGC AGCAGAAATC CAAAATTTGT ATGGGTAGAC CACAA

SEO ID NO:573: (Length of Sequence = 396 Nucleotides)

GAATCCCCAA AGGAGAGGA CTAACCTATG ACTATCAGTT TGATTTINAG GACGATCAGC ACAAGATCCC CTGCCACTGT
GGAGCCTGGA ATTGTCGGAA ATGGATGAAC TAAGAAGCTT TGAGGCTACC AGGCAGGGGA GTCCCCCTAC CCACAANCTC
TTCCCTGGAAA GNAAINGAGG GGGAAGAGAG GTAGCAGCCA GAGCCAGGAC CCAGGGTTGG GGCTGCCGGC TGACCCGGAG
CCCCTGGAGC AGGAGGCTGG GGCAGAGGGC CCTAGGCCAA GCCCACCCTG GGCACCAGGG ACAATCCTCT TCCCCACCAC
CGGCCCTCAG GCTGGCATCT CTGCCCCCAG CTTCCAGGAG GGGCCAGACA GAAGCAGCCA TTTGGCATCT CAGGTT

SEO ID NO:574: (Length of Sequence = 373 Nucleotides)

CTARACAGAT TIRACTCCCT CCCAGCARTC CAGATTARTT TRATATGCTT TCTTRACGGC ATTCCGCATT TMTCATTARA
GCARATGARC GICCATCCCT CTCTGATARA TTAGGGCARA ARARTTCATA TGTTTRAGGC ATAGGGRAGG AGGAGTTGTT
GGCTGTTARA ARARAGARCA ARARARAGTA CCGCARATGG CGTTTCARAGG TCTRGACATC TTCATCATCA ACACARACAT
TCCTCTTCAC ARAGGGRCCT CRAGTRACCT TRGGCTGGRG GRCCCACCTG CGTRTGTTTT TMTTCTCATT CTTTCTTTRC
CTTCCCTCCA GGCCACCCAA CCCACATTCA GTGGCCCAAG TCACGTGGGG TTT

SEO ID NO:575: (Length of Sequence = 431 Nucleotides)

GCCCCCATTA CCITCITIGC TGCTACCACA ACAAGGIATA TTAGCCCTTG AAATTAAAGA TGITGCTGTC CCAGTIGIGC
TTGTCTTCAC CTAAATGCAT ACAGTCATAT TCCAAAAGAC TATATATTAG TGATATCTAT ATAGTTCACC CTTCATATAC
ATGAGCTCCC GTGTGTGGAG TGAACTAATT GCAGATATAA AATATTTGGG AAAAAATTTC ATGTGTACTG AACATGTATA
GACTTTTTIN CTTGTTATCA TTTCCTAAAT AATACAGAAT AATAACCACT GTTTACATAG CATTTACATT GTGTTAGGTA
TTATAAAATAA TCTGTACATA ATTTAAACTG TACAGGAGAA TATGGCATAA GNCATATGTG GATACCACAC CATTTTATAT
CAAGTACTTG AGGCCTCTGC AGATTGTGGT G

SEO ID NO:576: (Length of Sequence = 410 Nucleotides)

GATGCAAACA GCCCCAAGGA GGGAGGTGGA AAGGCCAAGG GGCTTGCCCT CCTGCAAGCG CGCCTGTAAA CAAGTCCCCG
TGGGGTTTTG GGAGGTGCGC CCACATCTAA GACTGTGCGC CCTGCACTCC CTCTGGATGG CTTGCCGAAT TTGGTCTTCG
CTGATCACCA ATTCTGGAAG GGTGGAGAGA CAGTTGGCTG GACAGCTGCC TGATTCGGCC ATGACCCTTC ACGGGTGTCT
GTGGGCCAAC ACCAAACGCC AGCCTGCTCT GCTGGCAGGG CTTCTACCTG CACAGTCCCT AGGGCTGCAA GAGCAAATGG
GGACCCTGGC TNCCGGTCCT TNCCAGGGCC TTGGTCAATG ACATCACCAC TTTCTTAGGA CAGCGTCTTG GGGAGCTACC
GGAACTTTCG

SEO ID NO:577: (Length of Sequence = 405 Nucleotides)

GAATGAAAAT GGCATATTIG AACATAAACT TAGGGCAGAT TITTIACTACT TITIGAAAAAA TGITGGAAAA TATTICTGTA
TGAAACGTAA AACAACTITT AATTITTITT AGAAGTIGAG AGGATTCTAT TITIGCAAAGC TGTATTATGA AGCTAAAGAA
TATGATCTIG CTAAAAAAGTA AGTACAAACT GTAACATGTA TTCTTTTTTT AAAATCAATG CCTTINCTCA TTTNCTTCTT
TGAAATAGGT AAAAATATGT CCTTAGTAGT TCTTCCTAAG TGTATTCTGG AATAAGGGAT TTATCACTCA GACTGATGCT
AAGGACCAGC CTAGATTCCA TTGAGATTGA AACCGTAATT AGTGTTTTCT GCATGCTGCT GCTTTATACC AAGGGCAAGA
AATTG

CECTACAGGG GGGGCCTGAG GCACTGCAGA AAGTGGGCCT GAGCCTCGAG GATGACGGTG CTGCAGGAAC CCGTCCAGGC
TGCTATATGG CAAGCACTAA ACCACTATGC TTACCGAGAT GCGGTTTTCC TCGCAGAACG CCTTTATGCA GAAGTACACT
CAGAAGAAGC CTTGTTTTTA CTGGCAACCT GTTATTACCG CTCAGGAAAG GCATATAAAG CATATAGACT CTTGAAAGGA
CACAGTTGTA CTACACCGCA ATGCAAATAC CTGCTTGCAA AATGTTGTGT TGATCTCAGC AAGCTTGCAG AAGGGGAACA
AATCTTATCT GGTGGAGTGT TTAATAAGCA GAAAAGCCAT GATGATATTG TTACTGAGTT TGGTGATTCA GCTTGCTTTA
CTCTTT

SEQ ID NO:579: (Length of Sequence = 374 Nucleotides)

SEQ ID NO:578: (Length of Sequence = 406 Nucleotides)

GIGGGCCTGC TCTGGAGTCC ACATTCGTAA ATATTATGCT GCAGTAAATA TTAATCTTGA GAACTAGGTG ATATGGTTTG
GCTGTGGCCC ACTCAAATCT CATCTTGAAT TGTAGTTCCC ATAATCCCCA CATATCATGG GAGTAACCTG ATGGGAGGAA
ACTGAATCAT AGGGCAGTTA TTTCTATGCT GTCCTCATAA TAGTGAGTTT TCACTATATC TGCTGGTTTT ATAAGGGGCT
TTCCCCCCIN CCTTTGCTCT GCATTTCTCT TTCCGGCCAT TATGAGAGGA AGGACATGTT TGCTTCCCCT TCTGCCATGA
TTGTAAGTTT CCTGAGGCCT CTTGAGCCAT GCTGAACTGT GGAATTTAAT TAAA

SEO ID NO:580: (Length of Sequence = 396 Nucleotides)

CAGAATAAAC ATTTACIATT AGGAGAGICA AATCATTTAT TITCACATGA AAGAGATTAA GIAAAGCAGA ATCITTGATG
GTCIGCTGTG AATTCITCGC AGTGATTGAG AAATTCTGA AAACCACTTC CAAATCAATT ATAATATTAA GIAAACTTTG
GCTTTAGGAG TAAGAGAGAG AAGGTCTGCG TCCATGTTGG GAAAGAATAG ATATGCCCAC AATAATTAGT CTATTACTTG
TTTGAAAAAGG GTGATTTCCT CGTCATTTCA AAGTATTAAG CAAATAAGGA CATATTGAGT ATGIAATTCA TGGAAAANTA
AGNAACTTCT TACAGTATGA TTCCTAAAGG ATTATGGATG CCATTATCCA TTTTGGAGTT GGTATTGAAG CTTATC

SEO ID NO:581: (Length of Sequence = 449 Nucleotides)

SEO ID NO:582: (Lerigth of Sequence = 261 Nucleotides)

CCAGCAGGIC GIACTIGCAG TGGCAGGGIC CCGGACAGGG CCGCGTCAGI GIGCTGAGCI TGGTGGCGGG CACTGGCITC GACAGIGGCA TGACCCGAGG GAAGIGGCGG CGCGAGGGCC TCAGGGGGGCT GAGCACGICC TTGCAGAGGG GCGGGAACGG

GINCTGCTGG TAGTGGCCAA ANACCTCGAA AACAATGGGC TNGCTCTTGA TGTACAGGTG GCGTTTATTT TCATGGATTT ATACACACTG GAAAAGCCTC T

SEO ID NO:583: (Length of Sequence = 399 Nucleotides)

CCCAGGCCAC CATTIAAAGC AGCCATTCCT GCCAAAGAGC CAACATTGAG GCCAGCCGTT GCTCCAGCTA ATGTCTGCAG
GGCTCCAAGT GAGGCTATGG GGTTGACAGA ATTACTGCTG CTAGAGCTAG GTGAGGACCC TGAACTAGTG AGCACGCTGA
GGGGACTGCT GGATGTAGTG AGAGCATTGG TACCACTTGG TGTGTTCTGA NNTGCACTAG CTGCAGCAGC TAGTGCAGCN
AAATTCTGTA ACTGCATTGC ATTCAACCCT CCCATTGGGT GGAGGCTGCT CAGGGTGTTG AGGTTCCCAG AGGAGGCAGT
CTGCTGAAGG AGTGCTAAAT ACTNGGGTCC AAGAGTATTT AGACCAGCAA GGTTTCCCCA CACAGATGCT GCGCTGATT

SEO ID NO:584: (Length of Sequence = 441 Nucleotides)

GTIGITITIA AGGATIANAT GAGATATIC ATGTAATGTG CTCATCCCAG TGCCCAGCAC ACAAGAAATG TTCAATAAAA
TAGGAGGCAT AATTGTCCTG TITGAATACT AGATAACCCT TITAATGGAT ATTCTACAAT TATGAATCTA AGGTGCTTTG
GAGGAGCCTA GGCAATCTAT TCCAAAATTA AATGTAAGGA AGGTACATGA GTAAGGGATG GAGTAGGCCC TGGACCAACA
CTAGGAGCTCC AAATTTCCTA AAAAGCTTGA GCTTCTTTTA CTGTGGCCAC GCCTATAATG GGAATAAATC TGGTTCTTCA
AACAGTCCCT CCCCTCTCTA AGCTCTGCTG GGGAGTAGAG ACATCAGCAG GCTGGTTCTG TGNTTAGCTC CTCCCCATCT
TMGACTCTCA TCCCATTCCC TCTTTCCTAC TACCCATTCA G

<u>SEO ID NO:585:</u> (Length of Sequence = 326 Nucleotides)

GAAATGCAGG TTCAGCIAIT INCIGCTIGC AGAGTCCAGT TAACAAAAGT GAGINCTGT ATAAAGAAAG TNATTITITIT
TTITTAAATT ATTCCAAAGC TAGCTGAGGG GAACAAGTAC AGGCTTCCTG CCTAGGGGTA TCACTTTGCT TTTGGAGCAG
GAAGTAAGCA CTITTAAAGG GGGCTTAACA TGAATGGCAC ATGGGGTCGG GGGAAGTAAG CAAGTGCAGC ATCTACATGT
TAGTTTGGTA CCTTATCTAC TAGGTAGTCA AGGTGGTGAC TGCCTGINIC TTTGTGGGGC ATGTGTACTT TGGGGTTGTA
AATTGG

SEQ ID NO:586: (Length of Sequence = 431 Nucleotides)

GAACGAAGGA AAAGCATCAA AACCTACAGA GAAATINITC AAGAAAAAGA GCGGAGAGAG AGAGACTGC ATGAAGCATA
TAAGANCGCT CGGICCCAGG AGGAGGCAGA GGGGATCCTT CAACAGTACA TIGAGAGGTT CACCATCAGT GAGGCTGTTC
TCGAACGCTT GGAGATGCCA AAAATICTGG AAAGAAGCCA TICAACAGAG CCAAATITAT CCTCCTTCCT GAATGACCCC
AATCCCCATGA AATACCTGCG GCAACAGTCA CTGCCTCCAC CCAAATTCAC TGCCACTGTT GAAACCACCA TTGCTCGTGC
CAGINITCTG GGATACCAGC ATGTCAAGCA GGCAAGTGGG GTCINCAAGC AAAACTTGTC ACTTCCCAAA AGCAAGTGCC
TATGCTTGAC ANCCCAGGCC TTACTTCCCA G

SEO ID NO:587: (Length of Sequence = 338 Nucleotides)

CTCAAGCAAT TCTCCCACCT CAGCCTCCCA AATAGCTGGG ATCACTGGCA CAAACCACCA TGCCCAGCTA ATTTTGTATT
TTTTGTAGAG ACAGGGTTC ACCATGINGC CCAGGCTGGT CTCAACCTCC TGGGCTCAAG CAATCCTCCT GCCTCGGCCT
CCCAAAGTGC TGGGATTACA GATGTGAGCC ACCGCATCCA GCCCCACACC CTCATTTATA CCAATTACCT GCCCAGTAAC
TGTGGACTTT TGCTTCCTCA CCCCTGCTCT GATCTGGAAG GAGAGGGATT ATGTTATAGC TTGTCAGCAC AGTCCCAAAG
TTCAATATTT CTGCGGGC

SEO ID NO:588: (Length of Sequence = 277 Nucleotides)

AAGAACATTI AAGIAGITCA TACAAAGAAA TATAAATTGI NCITAAATAT ATCAAAATAT ACTCACCTCA TTCATAGTAA
AAGAAATAAA AAACTGIGCI CIGATGACAT TITTCATCTA TGAGATTTAC AAAGNICTAA AAATTGAGAA TATACATTTC
CIATTGCCTT TGGATGGCAA TTTGGCAGTA ACTATCAAAA GTATAAATAT CTATACCCTT TGAGGTGTCA ATCTCATTTT
AAAGAATTTA TTCTTCAGCT ATGTACATAC ATGTAGG

SEQ ID NO:589: (Length of Sequence = 353 Nucleotides)

GIAATGAATT ATAAGAATCT GAATTGAGAG CTAAAATATC TGGGTTGTAG GCCTACTCTG CCACGNITTT MITATTTGCA
AATATTAGAG CTGAACTAGA TGACCTCAAA GGCTCTAACC AACTCCAAAA CCTACAATTC AATGGCTGAC TGATATACAT
TGTATACTCT TTAAAAACAA TTAAAATCAA AGAMENTAAT AAATGTGTCA TGTATTATAC AACTATTATA CACGTGTGTG
TGTATATATA TATATNININ CACAGAGAGG AAAGACATCT ATACATAGNC ATAACCATCA AATCAGTCAG AATTTCCATC
AGACACTTIN CATTTCCCAG GTCCATCAGA TGG

SEO ID NO:590: (Length of Sequence = 364 Nucleotides)

CTCATATACA TAAAAAGTGA TAAGAATCOG AAAAGACAGC CAGGGGAATT AAATGCCAGT TGGGGCCAAC GGGGCCCTGA
TCACGGAAGA GGGCCCCCC AGCTCTCAAT CTTCACACAA TCCCTGCACC CAGGGTCACA GAGCATGCGC AGGTCCTTCC
CGCCCACTTC CGGGGCAACT GCCAACCACC GCGCAGGCTG AGCCCCAGGC AGGAAGCAGC CCACTTGGTG GGGTTGGGGT
ATGAGTCCTT CCTCGCGGGG GCTCGGTGGG TCCTGAGTAT TCTTTGGCCG GATTINCTGA TCCGTCTGCT CCAGGTGAGC
TNGGGAAGGC CCCAGGAAAA GGCCCANAAG GGCCTTTGCC AGGG

SEO ID NO:591: (Length of Sequence = 311 Nucleotides)

GAAAGGGGAA TAGGGAGTTA ACGITTAATC AATAGAGTTT GGGAAGATGA AAACGITCTA GAGATGAGTG GTGGTGATGC CACATAACAA TGTGAGGGTA CTTAATACCA CTGAACTGTA TGTTTAAAAT GGCAAAAAGG GTAAATTTTA TGTTATGTAT ATTTTACCAG AATTTTTTT TTAAAGCTTA CTGCATGGGG ACCAAGCGTG GTGGCTCACA CCTGTAATCC CAGCACTTTG GGAGGCCNAG GCGGGTGGGT CACTTGAGGT CAGGAGTTCG AGACCAGCCT AGCCAACATG TTGAAACCCC G

SEQ ID NO:592: (Length of Sequence = 358 Nucleotides)

ATTITISTITI CTACCCATCA TCCTCCTCTC AAAGGAACCA GOOGICCTIG GGGATTIGGC TGATGCCAGG GGATGGAGAG
TGICAGITIGG NTCTGAAGGG GAGGCTOGCA GCATGTGTGT GGCAGGTCAG ACAGACCCAA GAGCCAGCTT GGTGGGGCAT
CCCTGGCTAC CCTGGGGACA CAGTGAGCGC CGAACTAAAT AACATCAGGA ATGGNTCACA ACGCAATGAG TAAGGGGAAT
CTGAAGTCTAT AGGGATACAG ACCCAGAGGT AAATNGCCAT GGCCACCCAC TTTCCTACAG GAGAATGTGA CTAGTTGAGC
GTAGGAACAT GGGAACAAAT GGTAGAGGTG GCTGACAT

SEO ID NO:593: (Length of Sequence = 354 Mucleotides)

GACAGACTGA AGGAATATAT GCAGCTTAAT TTAACATTTT TTGAAATTTT ATATTGCAGA AGTTGTACAT ATTINCTGTT
GTGAAATTAG AAAGANTTGA CAGGCAAGGA GGGTGGTCTA CAAAGCACTC CATAGATCCA CCATACTGAG ACAATGCTTA
ATGCTTTGAT GGATTTATTT ATTINATACT TTCTATGCAT ATGCATGTAT TGTATAAATA CGNATGCATG GTTAAATAGA
AATGGTTCTC CTTGGTGTC TGTTTATCCA TTTATTGTTG TGAAGTAAAT CCCCAAAGAG GTAGGTTTGC TTTTGCCTGA
GGAGTCTTTT GCTACATACT GGCTGTACAT AATG

SEQ ID NO:594: (Length of Sequence = 319 Nucleotides)

GAACATGGCC GTGAACTGCT CGGAGATGCG CTTGAACAGC TCCTGGATGG CCGTGCTGTT CCCGATGAAG GTGGAGGACA
TCTTGAGGCC GCGGGGGGGG ATGTCACACA CGGCCACCTT CACGTTGTTG GGGATCCACT CCACGAAGTA GCTGCTGTTC
TTGCTCTGGA TGGCCAGCAT CTGCTCGTCC ACCTCCTTCA TGGACATGCG GCCCCGGAAC ACGGTGGCCA CCGTCAAGTA
GCGGCCGTGG CGCGGGTCGC AGGCGGCCAT CATGTTCTTG GCATCGAACA TCTGCTTGGG TGAGCTCGGG CACGGTCAA

SEO ID NO:595: (Length of Sequence = 370 Nucleotides)

GAAGAATANA AAGAAAAATC CAAAATGAAG AGTATTATAC AAGACAACTA GTCAATAGTC TTCAAAGTGT CAAGGTCATG
AAAAATTGAG GAAGCATCCC AGACTGAAGG GGACTAAAGA AAAGTGACAA CTAAATGTAA TGGGTGATTC TGGATTAGAT
CCTGGAATTG AAAAAGAACA TTCATGGAAC AACTGACAAA TTTGAATAAG GTCTGTAGAT CAGTAACAGT ATTGCATCAG
TGTTAATCTC CTGGTTTAGA TCATGTCCTA ATGGAAATGT TTTGTACTAT TTTTTGTGGA CTCTTAAGGA ATGTGGGTGG
AGGACACGGA TGAGACCTAC TTGCATCGAC AACAAGGCGT TCTACGGACA

SEO ID NO:596: (Length of Sequence = 335 Nucleotides)

CCACAGAGCC CCAACTCCCC CCACAGGAGC CAGCTCCCCC TCGAAGGCCT GGAGCAGCCG GCCTGTGACA CCTGAAGCCCG CCCATACAAG CCTGT
CCACTAGAGCC CCACAGGAGC CAGCTCCCCC TCGAAGGCCT GGAGCAGCCG GCCTGTGACA CCTGAAGCCCC CCATAGTTCTC
CCATACAAG CCTGT
CCCATACAAG CCTGT
CCCATACAAG CCTGT

SEO ID NO:597: (Length of Sequence = 336 Nucleotides)

CTCCTGAACA TCACAAACTT GGTTTCTACC TACCACACGA GTAGCCAAAA GAAAAGAAGC ACTAATAGAG AAAGGGGTGT CTCACACCAG ACAGAGGACC TCTGCTGTCA ATTAGATCCA GTATCATGAC CTAACTTTAA GTGTGGAAAA GAGTTCAGAT CTCTGAGACA CTGTGAAGAA ATGGATGGCT CATGTAACAT CTCTGATCCC TCAGTCCCCA ACCCTGGACG TGTTTCATTT ACAACATTCA TAGGAGTTAA CTTAGCAGTG TTGCAAGTTA AGGTTNCAAA CCAAATTATT TAATCAGTGT CCCCCCAATA AAATCACTTA TCCCATTTTA TTGCTAGTTT AGTTTT

SEO ID NO:598: (Length of Sequence = 402 Nucleotides)

SEO ID NO:599: (Length of Sequence = 369 Nucleotides)

CTCAACAAG TITGGATITI NICCACGATG ACTCCTTGGG TGAATITITA ATCAAGITAT TITCAACCATI TINCTCATAT
ATTTCGTGCA TCCCTATTCT GGTATTCAGT GAATACATGG GAGAGGTATG TNATTCTCAG CTCCCACAGC CCATAAGTCG
GGGAACCAGG ACTTCATTCC CCTCTGCTCT AACTCAGACT GTGAGGTCAT TGAGGGCGAG ACTGATGAAT TGITCCTCTT
CCTATCACTG GTGCCAAGCA CAGTAGTTGG CATAAAGAAG TTACTCAATA AAGAGGGGGT GAATTTAATG AAAGACAGAG
GAAGGAGGGAA CCTGGGGGAA GAGGTGGGCA TAAAGTGAAG GTACAAACA

SEO ID NO:600: (Length of Sequence = 342 Nucleocides)

COGCCTCCTG GGTTCAAGCA ATTCTCCTGC CTCAGCCTCC CGAGTAGCTG GGACTACAGG CGTGCGCTCC ACCACCACGC CCGGCTAATT TTTGTATTTT NAGTAAAGAT GGGGTTTCTC CATGTTGGCC AGGCTGGTCT TGAACTCCTG ACCTCAGGTC ATCCGCCCGC CTCGGCCTCC CAAAGTGCTG GGATTACAGG CGTGAGCACN CGCACCCGGC CAGCTGCTTC TATTTTAATC TGAACTTGGA AACACCTTCC TACTTTAAGG CACAGGATCA GGGTAAGAAC CCACATGTAC GAGCTAACAG AGCTGCACTT. CAAATTTACT TAAGTTAATT AA

SEO ID NO:601: (Length of Sequence = 319 Nucleotides)

AGIACTATIC IGCCATAAAA AAAAGAATGA GATCCIATCA CIIGCAACAT CIIGGAIGGA ACIGGAGGIC ATTATGIIAA GIGAAATAAG ICAGGCACAG AAAGAAAAAC IIIGCATATI CICACICATI IGIGAGAACI GAAAATIAAA ACAATIGANC ICACGGAAAT AGAGAGTATA AIGAIGGIII CCAGAGACIG GGAAAGGIAT IGGGIGGGG GCAGGGAATG GGGAAGGIIA ATAAGIACAA IGCAAIGAAT ACGAICINGI ATIITACAGC ACAAAAGGGI GGCIATGGIC AACAATAATI TATAGIACA

SEO ID NO:602: (Length of Sequence = 334 Nucleotides)

CACCCACAGA CIGCCAAGIG GGACAACITI CIGGCITITG AAAGGCICCI TCITCAGAGC ATIGGGGAGI CAGCAATGIC
CGITGIGITA AATCAGCIGC TGCCCATGAT TAAGCCIGIA ACCCAGAGAA CCAACGAGGA CIACAGCCCI GAGGAACIGC
TGATCCITCI CATATATATI TAINCIGICA CIGGAGAGCI CACGGIAGAC AAAGACCIGI GIGAAGCAGA AGAAAAAGIC
AAGAAAGCAT TGGCTCAGGI CITCTGIGAG GAATCIGGAT TGICACCITI GCIGCAAAAA ATIACGGACI GGGGACICTI
CAATTAATCI GACA

SEO ID NO:603: (Length of Sequence = 410 Nucleotides)

SEO ID NO:604: (Length of Sequence = 399 Nucleotides)

TCTCTAAGCA AAAAGAAAAT GATGAAAGAA GCAAACTTGG AGCATCAGAA AGGAAGAAAG AACATGATAA AATGAAAAATA
TGAGCTCCTA TIATGAACAT CGTATTACCA TTCATTGTGA AACTTAATCG TATATTTATA TATAAGCATC CTTCAGAGAT
GCTGTGGGTT CAGTTTCAGA CCACTACAAT AAAGTGAATA TAGCAATAAA GCAACTCATA TGAATTTTTT GGTTTCTCAG
TGCATATAAA ATTAANCTTC ATGCTATACT GTAGTCATTT AAGCATGCAA TAGCATTATG TCTAAAAANT GTACATACCT
TTATTTAAAA ACGCTTTTAT TGCTTAAAAN AGGCTAAATG GCCCATCTGA GCCATCGGCT TTTTTCCTGG CAGAGGGGG

SEO ID NO:605: (Length of Sequence = 372 Nucleotides)

ATGCCTTAGA AATCCTACCA CCTCCCAGAA ATGATAGITA TGGAAATTAA CATGGCATGT CAGATATGGT TCGCTGATGC
CTTGCTTTAG TTCTCAGAAA TAAGGCTTTA AAAGACTGGC ATGTTTCAGG ATTGCTGTCA GGAAATGATA ATTTAAAATA
CCCAAGAGTA CACTAAGAAT TATGGAAGCA TCTGTGAAAC TAATAAGCCA GTGGACATAC TGATTTTTAC CAATGTGTCT
ACATACTATA TTAAAAAACT TCCTACAAAG TATTGTCCCA ATTCAGTTCA TCTGAGGATG TGAAAACACT ACAGTGTACC
TTAAAAACATC ACATTCACAA CCCTGACAGA CTGAAATAAA ATGAAATTAG GG

. ..

SEQ ID NO:606: (Length of Sequence = 399 Nucleotides)

TECCTTCCTT TCTTCAATTC GAGACAGCAG TATCATTAGT GTTGTATAGG TTATAATTAA ATCTAAGTAG TTCTTTGTTA
AATCAAAGTT TACAGTAATA TCAAAGAAGA CTTGGCAAAC GTCAATAGTA TTCAGCAATT CACAAACATG GTCCTTAAAT
TCCATAACAT CTACAAATGT GAAGTAATAT AATGCCAGAT TTTNCAGAAT CTCTGATTTT CCTTTCTGTA GTTGTGCAAG
CTGTTGATTG TTGTTGCGGG TTTCTACAGC AGGGAATTTT CTGACTATGA ATTTCACAGC AGATTCCAGG NTTTTGTCGA
TAAGATAGGA TGGNTTTGCC NTGGGGCNCTC CACATGCCNT TCTTGATGTT GTAGAGGCGG GTGAGCATGC CGACGGCCC

SEO ID NO:607: (Length of Sequence = 412 Nucleotides)

CTGTACCCTT ATAAAGAGIG AAAGCCCTGC CCCCTTCTCC TATAGAACCC CTAGCAAGGA GACTGGAAGA NTCAAAAACA
ATCCACCCAA AAAATGGCCT GCAGGGACAC AGTCCCAGAG AAAGAGACTA TGTACAACAA GGTACAGTAA GTAAGACCTG
CCCACACACA GGACTTCCAA TCGACTTCTT AGTGCTTACT CCTACAGATG AACAGATCAA CCAGGGCCAC CAGATGCTCC
AGGAAAGACA GGAGTCCAAA AAGAAAATTC GGTAAGTTTG AATATATTTT GAGCAAATTT TCAGTTCTGT TGAAGTATTG
GGGGGACATT CAACAGTGAG TAGTAGTTTA GGGGGAACAG CTGGCACCTC TGGCAGTCGC CTCAGAGGTC AANCCAGCGT
NTAGGTTGCT TT

SEO ID NO:608: (Length of Sequence = 419 Nucleotides)

ATGAAGGCAG CTGAACTCTC CATCAAGTTT CTGCCTCCCC AACGTAATAT GGAAGTCGTT CTGGCTGTAG GACCCCAGCT
GATTGGAATT GGAAAGCACA GTGCAGCTGC AGAGCTCTAT CTGAATCTGG ACCTTGTCAA GGAAGCAATC GATGCTTTCA
TCGAGGGTGA GGAGTGGAAC AAGGCGAAGG TTGTAGCTTAA GGAGTTAGAT CCCAGGTATG AAGACTATGT GGACCAGCAT
TATAAAGAGT TCCTCAAGAA TCAGGGCAAA GTGGACTCGC TGGTGGGTGT GGATGTGATA GCTGCTTTGG ACCTGTATGT
GGAGCAGGGC CAGTGGGGAC AAGTGCATTG AAACAGCTAC CAAGCAGAAC TACAAGATTC TGCACAAGTA TGTGGCTTTG
TATGCAACTC ACTTGATCC

SEO ID NO:609: (Length of Sequence = 337 Nucleotides)

GGIGGAAGIT GIAGIGAGCC GAGATCATGC CACTGCACTC CAGGITGGGI GACAGAGAA GGCTCCATCI CATAAAAAAA GAAAAGAAAA AGCATTICIG AAAGGAATAA AAAACAAATT GATAACATCC CCTAATCICT AGITGITGGG ATGIAGIATC CTTCATTIGA TCAGGAAATC ATATGATTGI CCTTAAATTA TTAAGITGGC AGAATTIGIG TGGTTTCATA ATGATGCTIG TAAGATGATA TINTAATGGA AATGITTTAG ACTATATCIN TIGINGITIT TNCTGCTGIN TITGIGIAAG GCTTAAANCT ACCCCCTTTA AAAACAG

SEO ID NO:610: (Length of Sequence = 441 Nucleotides)

TAAGCCAGAG ACATTICACT GIATTAATCT TGATACTAAT TACTAAGGCT TITCIGIGGA CATTAAATTI GATCIGITTA
ATTGCAAATA CAATAAAAGI CGIGATTIAT GCITAATGIT TCIGCTAGGC TGATGACATT TIGAAAATAG CACTTATAGC
CIGGTTTGIC TIGGTTACAA CITTIGIGGC TCCAGATGCT AAAAAAAATC TAATTGAGTA AGTAAATAAT GCAGCTAAGC
GIGCCICTCT CGCITCCGAA AAGITTTTIC TACTCCTITT TCTCCCTGGA GAGGCCCTGC TGCACACTGA TGCTGATCTA
AGGAAATGCC TITGCTTCTT TGCCACTGAG CAATGITAGA ATCACTAGGA GGGCAGGGCT ATCCCACTGG TCACTCTGTC
CCAGCATATC TACCATGAAG TCAGCAGGGA CTACAAACTC C

SEO ID NO:611: (Length of Sequence = 344 Nucleotides)

 GIACATATAT GIGIATATAT GIATATATCC CACATCICCA ATTINOCIAT ACGIATATAC ACACATATAT GITATATAGG GIGIACAGAT ATAGGATATG TOTG

SEO ID NO:612: (Length of Sequence = 384 Nucleotides)

TGATGACCAT AAGCCCATGC TITICATAGA TGITTAAGGG TIAAATGAGG TAATGCATGI CGAGTGCTCA GCCAACTGAG
ATTCAGGAAG CGCTCAATAG ATGCTGGCTG TCATTATTAA CTGAGTAAGT AATCCTTTTC CCACAGAAGC AGTAGAAGGC
TGACGATGTG TGTGAAAAGG ATGGATACAA TICCCTGGGC CACAAATAAA GGITTTTTTG GTTGTTGTTG TTGTTTAAAT
GAACTGAAAT GAGTTTGAGA GATTCATATA TIATTTTACA ATACTTCTTA ATGCTAGTTT AAAAAGTTCA ACATTGTCAT
TCTACTCCAC TTCCGTATGA GATAAGTATA TGAGGGNGCT TAATTCCCCG MTAAACTAAG CAAG

SEO ID NO:613: (Length of Sequence = 342 Nucleotides)

TATTIATITI TGIGGGIGIC GACITCCIAT GIGGGCITIT TGGGIGACAC TCCCTIAAGG GITCAGITIG ACAATICINA
GAGITGICCI GCAGITGGAG GCCACCAGAG GIATCIAAGC TCCCIGCTIC CIATTINATA ATCCICCAGC CCCAGCAGGI
CCACICCIGG TICCIGIGIG TITGGCCCGG GCACAATCCC CACIGCTITG CIAGACGIGC TITCIGCCAT GIGGCITIGG
GCCIAGAGCT TGITGATAAT TGCAGCTIGI GGCAGIGGAA ATATGGCIGA ATGAGCGICT AAACCCCIGG GINGGGGCNC
TNAANINCNN GGGTTTTTAA AA

SEO ID NO:614: (Length of Sequence = 393 Nucleotides)

CAGIGITATI AACAATAGCC AGGAGGIGGA AGCCACCTAA ATGICCATCA ACAGATGGAT GGATAAATGA AATGIGGICT
ATACATACAA TGGAATATTA TICAGCTITA AAAAAGGAGC AAATCCTGCC ATGIGCTACA ACGIGGATGA ACCITGAGGA
TGITTIGCTA AGIGACATAA GCCAGICACA AAAAGACAAA CGCTGCATGA TICCATITAT ATGAGGAATC TAAAGITAGIC
AAACTCTTAG AAAGIAGAAT AGIGGITAGC AGGGGITAGG GGGAGGGGAA AAAGAAAAGT TACTGITTAA TGGCTATAGA
GTITCAGATA TGCAATACGN NAATTTCTGG GGGATTCTIT TGCACCACCA ATGIGCACCG TATAATTCCA CTT

SEO ID NO:615: (Length of Sequence = 310 Nucleotides)

ATTATATACA TICCITTACT GATITITAA AATTGIGICA ATATCITCAG TGAACICITA ACAATCIGGG GAACIGITIT
CCICAATTAC CACTICAGCA ACGITCATAC GAAATCAAGG CITGCCITCA TGICAGIGIC AGGITCAACT TTAACTCGAA
GGITTIGIGIT TGICICTAAC ATCITCAGAG TGAGCITTAG GGATGCCTGA AGGATGGACA GIACAAGCAA GCAGCTACTT
CCATGATACA GIGGGAAGAT AAAAAGGCCC ATTCAGICCA GCCGIGACCT GIAAATCCAG CTIGCCCTCC

SEO ID NO:616: (Length of Sequence = 266 Nucleotides)

GAGATGGAGT CTCGCTCTAT CACCCAGGCT GGAGTTCAGT GGCACGATCT CGACTCACTG CAAGCNCCGC CCCCCAGGTT CACGCCCATIN TCCTGCCTCA NCCTCTGGAG CAGCTGGGAC TACTGGTGCC CACCACCACT CCCAGCTAAT TTTTTINTATT TTTGGTAGAG ACGCGGTTTC ACGGTGTTAG CCAGGATGGT CTCGATCTCC TGACCTCGTG ATCCACCCGC NTCGGGCTCC CAAAAGTGCTG GGATTACGAG CGTAAG

SEO ID NO:617: (Length of Sequence = 376 Nucleotides)

ATAATAATGA AAAGTGAAGG GTGGGGGTGC TGGCCACCTC CCATTTCTTT GCCTGGGTGG TGGTGACCAC GGCGCCCTTG
TGTCCTTTCC ATTGGTTACT GAGGACCATT GCCCTCATGG GCCAGGCCA CAGGCACCCA CCTGTNAGCC TCACCTGCCA
CCTCTCTCCA TGTTGGCTIN TTGCCCCTGG GGCTGGCCTG GGCATGGGGG AGCTTAINIC CCCGACCAGG GGCTTGGCCA
TGINICCTTC ACAANCCCCA CTCCCCGGG ACTGAGCCTC CACTCTCTGC TGGGCTGAGG GCTCTGTGT NGCCCAGGAG
CCCTCCCAGC CACGTGCCAG CCCATCCCAT CATCAGCACT TGGTTTTAAG CTTCAA

SEQ ID NO:618: (Length of Sequence = 352 Nucleotides)

GCCCATCCTG GCTAACACGG TGAACCCCGT CTCTACTAAA AATACAAAAA ATTAGCCAGG CGTGGTGGCG GGTGCCTGTA
GTCCCAGCTA CTTGGGAGGC TGAGGCAGGA GAATGGCATG AACCCGGGAG GTGGAGCTTG CAATGAGCCA AGACTGCGCC
ACTGCACTCC AGCATGGGC ACGGAGCAAG ACTCTGTCTC AAAAAAAATAA TAATAATAAT AAAATAAAAA GTTTGTTAGT
ATTAGCAGAT ACATATTACT AGGTACCCCC CATGCTCAAT GAAGTGTTGG GVTACTCTNA AAAAGTGTCC AATCTTACAG
GTGTGACTTC CTCTGGAACT GCAAATTCTT TT

SEQ ID NO:619: (Length of Sequence = 359 Nucleotides)

AAAAAAAACG ACCCCCACAA GGGGGAAGGC CCCAAGTGGG CCCCTGCCTG TNGTNCTCTC TGGCTCCAGA GATGTCTGCA
TAGGCCTCAG CTTCTCACTG GCCAATCTCC TCTTCATGGG CACCAGCCAC TGCTAAACAT CCTTCCCTCA CTTCTTGTGT
AAGCTTGCTC CCCTGAGCCA CAGGTTGCAC ATCTAAACCT CAGCTCCAGG GAAAGGAAGA ACCAATGGAA GTGCCAGAGT
CCTGGGGCAA GCCAGAGCAT CACCTGTCAG CAAACCTCTG CTGGGCACTC TAAGCAAGCA CAGGACAAGN CCCAGAGTTT
AGTGTGTCCA GTATCCAGCA TGGNGACAGC ACATGCATT

SEO ID NO:620: (Length of Sequence = 447 Nucleotides)

CTCTCTCAGC ACAGCCTGGG GAGGGGGTCA TIGITCTCCT CGTCCATCAG GGATCTCAGA GGCTCAGAGA CTGCAAGCTG
CTTGCCCAAG TCACACAGCT AGTGAAGACC AGAGCAGTTT CATCTGGTTG TGACTCTAAG CTCAGTGGTC TCTCCACTAC
CCCACACCAG CCTTGGTGCC ACCAAAAGTG CTCCCCAAAA GGAAGGAGAA TGGCAGCCTC CACATCTCGG GTTCAAGTGA
TTATCCTGCC TCAGCCTCCA AGTAGCTGGG ATTGCAGGTG TGCACCACCA TGCCTGGGAT AATTTTTTGT ATTTTTTAAG
TAGGACACGG TTTCACCATG TTTGGCCAGG CTGGTCTTGG AACTTCTTGA GGTGTAAATG ATCTTNCCCC ACCTTNTGCC
TTCCCAAGTG CTTGGGATTT ACAAGGTTTT AAGACCACCCG AATCCAT

SEO ID NO:621: (Length of Sequence = 237 Nucleotides)

CAATACCCCT GENTCCTGGG GCAGGTGTTC TGGGATCCTG GACAGGAGGG TCAGGTCGAT TTTAACCCAG AGAGACCTGA
TCTCATCACT GTCCTTTAGA GGGGAGAGA GTTCGTNCCG GCCAAAGGGG ACCAGTGTGT AGAACTGCTC CTCCAGCTCC
TTGGCGATGT CACTMGTGGT CCTGGCGTTN ATGGAGCCTA CAGGGGCCCT AGGACCACTG CCCCCMTTGG CAGCGGC

SEO ID NO:622: (Length of Sequence = 247 Nucleotides)

AGAAGGICAA TAATAACAAA CITCITCAAG GIAAAGCAGG AIGITGGAAA CCATIGCAAG GAAGCTAAAA ACCITGAAAA AAGATTAGAA GAATGGCTAA CIAGAATAAA CAGIGTAGAG AAGACCTTAA AIGACCIGAT GGAGCIGAAA ACCATGGCAC GAGAACTACG TGATGCATGC ACAAGCTICA ATAGACAATT CGATCAAGIG GAAGAAAGGG TATCAGIGAT TGAAGATCAA ATAAATG

SEO ID NO:623: (Length of Sequence = 315 Nucleotides)

AATTTAGGIT TGTTTTATTT AAGITTAATG TTAATTCCAT GCTGTGTTTC AGTAAGANCA ATACAGATTC TGTATCTGTG
GCTCCAGTCA GATATCCAGT AGTACAAATN AGCTTCAAGT TACACATACT GANCAAAAGA GGTTGAGCGA GCGAAGGAGG
GGAGGAGTGA GGGGAAGGAG GTAGGGGGAG GGGGAAGGAG AAGAAACAAA AGANTTGAAC AGGCATGCAG GCTTTTCCAT
ACCACCTTCA ACGCTAACCT GCTTCAGTGG GAGAGTAAAG TAGGCAAGAN TGAGCAGCCA CGGATTGTTG AACTG

SEQ ID NO:624: (Length of Sequence = 375 Nucleotides)

 SEO ID NO:625: (Length of Sequence = 305 Nucleotides)

GITCCIAGAT TACICAAATT TAGIACICIT CCATCITTIC TIGITGCIAT TCITITAAAA TCACAAGAAG TCCATAACIT
AAGIAGGAAT TIGIATAATG TAACITATIG TGAGIATATT TCCITACCAG CTCATAAAGA ACTATGIAAA CITGAATGCA
TATTITINAC ATAAAAATAG CAAAAAAAAA AAAANCAAAA AAAAAACAGI ACTGGCCTAA TACTAGINGA NITACAGAAT
ANGGGIAAAAT ANTACATGNN CATCCITACA GAGIGAGCAT AAACAATACA TGGTAATAAT ATTTA

SEO ID NO:626: (Length of Sequence = 300 Nucleotides)

AGCAATCACA TAAGGAAGGC ACCTOGAGIC TAGTAACACT GTGACTCTTG CGGTCTCTTA GAGGTACTTG GTGGTCTTGG ATAAGATCTG GAAGAATTCT TTGGATTTCC AGACATAGGC TCTTGTNCTC TTCCCCTAAC TTCTCCCCAAA CAAATGGCAT CTCTCTCTCT CTCTCTGT GCTGAGCTGC CTAGAACTGT GGGTGGGATC ACACAAGCAC CCTTNTGGCC ATTGCCCTTG GGACTGTGCT AGGTCAGACC TGAAGTCAGC ACAGCATTGG GTCTCACCCA ACACCTGTGG

SEO ID NO:627: (Length of Sequence = 369 Nucleotides)

GAAAAAGAGA GAGGAGAGG AGTCAGGAGT GCTTTGGAAC TGGAGGTTTG CTTTCCACTG ACAACATCCA TATCINCTGC
TAATGCCAAC ATGCTCCCAA GTGTCTTAGT GGGTCCCACA AAGTTGATCC AGCCCAGAAG AGTTGCAGGG ACAGTCAAGA
AACCAGAGGT GCTGCCCACA TCCCCATCAC TCCCTTTCCC AACTTCCCAG CCTTGCCCCA AAAGCAGCAG CTCAGGACAA
CCTGAGATAC TACTGTNATG GGTCCCCGGG AGGAGGACAG CAGGAGTCTG AACTCCAGAG GAGGGGGAAT ATGGGTAAAA
CAGAGAGATG GCAAGGAGAC AAGCTGTNCC CAGACAGAGG GATGGGAGG

SEO ID NO:628: (Length of Sequence = 310 Nucleotides)

TTTTTTTTT TGAGACAAGA GICTCACTCI ATCACCCAGG CIGGAGIGCA GIGACATAAT CATGGCTCAA TGCAGCCTCG ACCICCAGA CICAAGIGAT CCICCCACCT CAACATCCCA AGIAGCTGGG ACTACAGGAG AGCCACCATG CCCAGCTAGT TTTTNACTTT TCIGCAGAGA TGGIGITTCI CCATGITGCC CAGGICGGIC TCGGAACTCC GGGGCTCCAG CGATCCTCCT GCCCCAGAGTC CCCAGAGTGC TGGACCCACA GGCATGAGCC ACCACACTCA GCCCCCAAAAT CCATGATTTT

SEO ID NO:629: (Length of Sequence = 443 Nucleotides)

CGCAGAGCAG AGGGGAAA GGCAAAGAGI ACAAGIGAGC GAGCCCTTT TGIGATGGCG TIGATCIGIT TACAAGGGGA
CTGCCIAAAC ACTITCCATT AGCCCCACT TCCCAACACT GITGCAGIGT TGCAGITAAG TITCCAACAC ATGAATGCTG
GGGGACACAT TIAAATTAGA GCAGIGATGA TCAGAAAGIT ATTGITGGGA AAGGAGGITC TATTITAACT TAAGIAGCTT
GAAAAAGCTC TTCAAGGAGI TGATACAAGA ACTGAGATTT GAATTAGAGG ACCGAGTAAA GTGAAGAATC TGCGGGCAAA
GTCCCAGGCA GAGGGAAGAG CAGGAAATGA TTCATCAGTA GACTTGCTCT CCCATTCTCG GCAAGGGCTA TTTCACATTT
TCTTCCACTC TCTTCCTCAG CACATCTCCA CCTGGGTTTT CTC

SEQ ID NO:630: (Length of Sequence = 263 Nucleotides)

TOGATGREET GAAAAGCGAA CACTIATAGA CIGCTACIGG GAACGTAAGT NAGTACAACC TCTATGGAAA ACTGTATGGA
GATTTTTTAA AGAACTAAAA GTATATCTAC CATTTGATCC AGCAATCCCA CIGCTGGGTA TCTACTCAAA GGAAAATAAG

TCATTACATC AAAAACACAC CTGCACACAT ATNITTATTG CAACACAATT CACAATTGTA AAGATATGGA ACCAACCTAA GTGCCCATCA ACCCAATGTA GGG

SEO ID NO:631: (Length of Sequence = 221 Nucleotides)

AATTTINACA TATCAGIAAT TGITTITATA ATTTGTGGIT TINATGAAAC ATTGCTATGC ATTTATTAGG AAAAACTGAA
TTTCCCAACA GGTGAACTGA AAAGNTATTT TAACTATTAT ACATAATCAA GATCCTGCCT CTACGGAATT AGCTAAACCT
AAAAATGTTT GCATTAATGN ATAAATTCTT CCNGCATTCC TTGGGCCNGN TCTGGAAGTG G

SEO ID NO:632: (Length of Sequence = 344 Nucleotides)

TGTGATGGAG ACAAATACTT CAGTATTGGG ACCCATGGGA GGTGGTCTCA CCCTTACCAC AGGACTAAAT CCAAGCTTGC
CAACTTCTCA ATCTTTGTNC CCTTCTGCTA GCAAAGGATT GCTACCCATG TNTCATCACC AGGACTTACA TTCCTTCCCT
GCAGCTACTC AAAGTAGTTT CCCACCAAAC ATCAGCAATC CTCCTTCAGG CCTGCTTATT GGGGTTCAGC CTCCTCCGGN
TCCCCAACTT TTGGTTTCAG AATCCAGCCA GAGGACAGAC CTCAGTACCA CAGTAGCCAC TCCATCCTCT GGACTCAAGA
AAAGACCCAT ATCTCGTCTA CAGA

SEO ID NO:633: (Length of Sequence = 378 Nucleotides)

GGTCAGACCT GAAGCCGGCA CAGGCTGTG ACTGCCCAAG ACCCCCACTG TAACAACAAC CCAGCTGCCA CCTATTTCAC
TCAAGGCCCC AGGGCTCTCC AATTAGCAGG TAGTGAAGCC AGCCAGGCTT CTNTCCTTCC CTTCAGTGCA GTAAGCTCCC
CTGGTCCCTA GATGCATTCA AAGGTGCTGT CTGAGAGCCA GGGCTCTCAG TCATAAACCT TATAAATCTA CCTGGNGTTC
TGTTCTACCA TCGCTGAGCT GGCACTGAAT CCACCCGGCA AATCCCTTCC CACTNTCCCC TCCCCTCTTN CCCAGGCAGG
GTAGTCTGTT NCCACCTACG ACGTCATCAC AGTCTCATGC GGGATTACTG CCAGCTTC

SEO ID NO:634: (Length of Sequence = 28 Nucleotides)

ATCAGTGGTC TACCACAGNT TAAGTAACGG GTCATATT: G GAGTATCACA CATCTCAGTC TTGTAGAAAT TAGCNACAGC
AATTAGGAGT CATGCACATA TANGAGATGT AATCCCACCC TTTGACTATA GCCTACTCTT GINTITTACA GAAAAGACTG
TGGNGGAAGA AAACCCTTTA CCCINTINIT CAGGGAGAAA CTNACANCAC TCANCTGCCT GGCACTGAAA ATNTGGCATC
CAGTCCACTT TACCATCAGT GITTAAGGAA ACCATCTCTG GTAAGC

SEO ID NO:635: (Length of Sequence = 226 Nucleotides)

TTGGGATGAT GCTTTTATTA AACGGAAGCG TCCAAAAAGG TCTGAGTCAA TGGTGGAGAG GGCAGTCAGC CCTGTGGCAT
TTCAGGGCTC CCCACCGATA GINATCGGCA GTGCTNACTG CAATGTGATA GAGATAGATG ATACCCTCGA CGACTCCGAT
GAAGGATGTG ATCCTGGTGG AGTCTCAGGA CCCTCCACTT CCATCCTGGG NGTGCCCCTC CCCTCA

SEO ID NO:636: (Length of Sequence = 367 Nucleotides)

AACGCAATAA AAAGACAAAT TOCAAAATGG GCAAAAGATC TGAATAAACA TITCTOCAAA GATATGCAAA CAGCCAATAA ATACATGAAA AGATGGCCAA CATCATTCAT TATGCATTGC AGAAATGTAA GTCAAAAACCA CAATGACATA CCACGTTGCT CCCACTAGGA TAGCTACAAT CAACAAAATG GACAGCAAAA AGTGTTGGTG AGGAGTAGAG AAATCTGAAC CCTCATGTAT TGCTAATGGA AACACAAAAT GATGGAGCTA CCATGAAAAA CTGCTTATCA GTTTGACCTC GGGAAGTTAA ACACAGAAGT ACCACATGAT CCAGCAATTC CACTCCTAGG TATATACCCC AAGGACT

SEO ID NO:637: (Length of Sequence - 384 Nucleotides)

TTCATAAAAA TITTACTTAA AATCIGTAAC GCIAGATATT GACTATCCTT AGTIGAGICA CIGAGGITTA AACACAATGG
TAAGICTTAA AGICIGCTAT TTACAGAGCA TIGAATCIGT ACCAATTIGC AATAGAAAGC CITCAGTATG CAAGAAGTIT
GCATGGGIAT TAAGAACACA GCCIAAATAA GGCATTIGAT CIAATCIGCA GGAAGAATTT TCTTCCCCCAA AACAGAATTA
TAAAAGCTTA CITTAAACAG GAGGCAGAAT AATTCTTTTA GGAAACCATT TCATTCTGIT TCTACTAACC TATACCATCT
GAGGAATTCT AGGGAGGATA ATAAAANTCT CGTGTATTCC ACAGCAAACT TACATACCCT AAAG

SEO ID NO:638: (Length of Sequence = 409 Nucleotides)

GAAATTITIC AICAGCICTI GITICCICIC ATTCITTITG ACCTIGIAGA TITATCCITI TITCITAAIT TATTCICACT
TAATGGGATI TCAGGAGCAT ATTGACIAAG TITTCATTIT TACATGIATA CIGGGGAGIA TGACATAGAC ATCICTGIAC
TTAGATATIA CIGATGIAAG TCTACITTGA ATCAAATGAA CAGATGITTA AAAAGIATTG TNOCCAATTG TITTTAATGAT
TTCINCCIGI GAGTIGGGGI GGIGCIGCCC ATCACCAACT CAGGACGGGI ATTTGAAAAT ACCTGGGNNA AATTGIAACA
ATGICTGGGA AAACACTGCA GGATATITTA ATTGGGCAGA GGGGTCAAGG GGATGGATTA ACCATTGCGG AAATGIGAGG
GACGGGTCC

SEQ ID NO:639: (Length of Sequence = 197 Nucleotides)

GGITCIACIC ACGGCICAAG AGCATGGCIC AGGAGGAGAI CCGCAGAGAG AIGGACAAGA INATCGAGGA CCIGGAGCIC ICCAACAAAC GGCACTCACI GGIGCAGACA ITGICGGGIG GCAIGAAGAG CAAGIGIACC GIGGCCAIGG CCIITCGIGGG CGGCCICTCGC GCCATCAICC IGGACGAGCC CACGGCG

SEO ID NO:640: (Length of Sequence = 398 Nucleotides)

GAGAAGGAGT TICGCICTIG TCACCCAGGC IGGAGTGCAA IGGIGCGGGC ICGGCICACT GCAACCICTIG CCICCCCCGG GITCAAGGGA TICTCCIGCC ICAGCCICCI GAGGAGCIGG GATTACAGGC ACCGGCCACA CACCCAGCIA ATTITCIATI TCCAGIAGAG AIGGGGITTC ACCAIGITGG CCAGGCIGGT ITTGAACTCC IGACCICAGT IGATCIGCCI GCCICGGCCT CCCAAAGTGC IGGGATTACA GGCGIGAGCC AITGGCACAC AGCCITATCI GCATTITCAA ACGGGCCAGI AIGGATGGGT ITTACACTIA IACINGAAAG GICATCCITT INAAAAAANG AACCITTAAA ACCATTAACT ATATATAAAA ACTATATT

SEO ID NO:641: (Length of Sequence = 402 Nucleotides)

ATAATTIINA GCAAAATGAT ACAAAACINI NITAACCAAG TAGAAGATIG GIAGITACAG TAGAATCGIC AGGGAGTACA
GGGCGGCCAC CACTGGAGGG AGCTGAGGCC CTGGAAAAGG AGTCTGATTC TNIGCAATTC TCTCTCTGCT TTINITCCCA
GCCCCGITAC AACCGAGTTC ACGTGGGGGG CCGCAGTGCA GCCCCAGCGG TGGCAGCTCT TGGAGTCTGT CCGTTTAGTA
TGTTTCCCCC ACGAGCGTCG CTGGGTGAGT GGCCTGGAGA GCTCCCGGTG TTAACATTTC GATCCTAGAC CCGGGGGACG
TGTCACTAGG TAAAGGCCAT TGGGTAACCA GAGTAGATCA GGCCATGGCA TTTGTCTGGC CCCTTTCACA GCAATTAAGG
GG

SEQ ID NO:642: (Length of Sequence = 395 Nucleotides)

SEO ID NO:643: (Length of Sequence = 325 Nucleotides)

GGTATCTTAA AGCCTTTCAG GGATTTCAAT AGACACATTT CTTTAGCTGA AATCTATTCT CTCAGAAACT TACCCAAACT
TCTTAATAAT GINCAAATTC TAAGAAAGAT ATCATGGCTA CACAGCACCA GGNAGAGCAC ATTATTTCTC TTCACAATTC
CCTTGCATAG CATCATGGCT TCCTAAGGGC TTTTAAGTTT ATTGCTTCAA CTGATTCTCA TAAAATCTCT GAGATGCTAT
CTGGAAAGTA TTATTATCCC CAGTTTGCAG ATAAGGCAAC TGAGGTCTAG ACTTGCTAAA AAATCACACA ACCAGGTAAG
TGGGC

SEO ID NO:644: (Length of Sequence = 373 Nucleotides)

CTTCACATCA GCAGCCGGAC GAGGIGACTG AAAATCCAAA ACAGAAAATT GCAGCAGAAA GCAGTGAAAA TGTTGATTGT
CCAGAGAATC CTAAAATGAA GTTGGATGGA AAACTTGACC AAGAAGGCAA TGATGTAAAA ACAGCAGCTG AGGAGGTACT
AGCTGGTAGA GACACATTAG ATTTTGAGGA TGTCACAGTT CAATCATCAG GCCCGAGGGC TGGTGGTGAA GAATTAGATG
AAAGGTGTTGC AAAAGATAAT GCTAAAATAG ATGGTGCCAC TTTAAAGCAA TCCTNGAAGG ANCCAGAGGA GCGAAGGATG
CAGATCACTG CACCCGTACC CCAAAAATTG GAAAGTCCCC TCACAGGCCA TTT

SEO ID NO:645: (Length of Sequence = 310 Nucleotides)

TTTTTTTTT AAGACTCAAG GTAATGAAAA CTATGAGTAG AATAGTAAGG TGTGACAGGG GACAAATAAG TAGATATAAA ACTATGCTGC AATATTTTAG TTATTAAAGC TGGGAAATAT GCAAATGTAA GTAGTGCTTG GAACCAGAGA AGGTTCTATA TTTAGCTGTT CTTCTGTAGC TAAATCTGAC AAATTGAAAA ATATCATATT CTCTGCTCTA GGTACATTTT ATGTATATTT TGACAGCATA TCAAATATAT GANACATTAG GTTAAATAAA TTAAAATCCA GTGGGATAAA CTATATGOGG

SEO ID NO:646: (Length of Sequence = 362 Nucleotides)

CTTGGGATTG CTAGATCAGT GITTTAGACA GGAATGCCAA GGCAGAAAAG AATCACATAT CCAGGACCAC ATAAAANCTG
GAGTGTATGT CATAACAAAT TINCTCCTGT GCTTAGAAGT TITATGGCTT TGGATTTTAC ATTGATGTTT GCAGTCCATT
TTGAGTTACT TTTTGTATCT GATATGAAAT ATACCCAAGT NCATTTAAAA AATAAGATTA TACAGTTGTT TATGGAATGC
ATTTATGTAC ACGGGTAATC TGTTTTGATT TTGTGTGTAT GTTAAAACAT CTTTATTATA GTATTNTGTA AGAGTAGGTT
AATATTGACC TTGGGCATTT TTAAACCAAG GGGGGAATTT CC

SEQ ID NO:647: (Length of Sequence = 226 Nucleotides)

TTTTGGCGTC AGATCTGTAA GTTTATTTGC TCAATGTACG ACAGCTACAT AATGNCTTAC ATTCATGATA TTCCATCACT
GAGGAAACTG CTAAAGATGG TCCGTGTGG AAATAATTCC TTAGAGAAAC ACGGAGCTGG AAAAATAATC ACTGATTAGA
CCTTAAAAAAT AGTTCACTGC ATAACATGNC AAAAAGCACA AAGGCTCATT CAGAGAACAT ATTTGT

SEO ID NO:648: (Length of Sequence = 198 Nucleotides)

AACTAAAAAG TTAAAACTIT TACAAAACAA CAAGIITITCC TTAAATTATG ATTTGITATT ATAAAANCTA GTAAGAAAAA ATTCCACCAC ATGAAAGCAT TINCIAAAAT TCATACCCCC GTACCTATIT TTAANTACAG TTGGTAAATT GATTAAGCTC TATTINCATT TIGANIGATC ATCGGTTTTA TTTTATTT

SEO ID NO:649: (Length of Sequence = 337 Nucleotides)

ACATCIGCAG CCATATATGA GGICCCTCAT GAGACTIAGC AACAAGGIGT GTTTTAATGI GACAGIGIGT CTGATGIGIC CCCAGCACAT TGGGACCAGT ACACAGIGIT ATTIGTACAT CIGCIGAGTA ACATTGAGIG TGIGGGTAAC TAAAGCCCTC AGTAATTATT TTACITAATG TTTTCAAGCI TAATTCIGAT CTTGTACTIG CATGATTTAT TATTCCTTGI GCTAAATTCT TCAATGITCT TGCCTTGATT GATCTGCAT TATCTATCAC TTAACTAAAA TANTAAATNC CTTTAATTAA GICATGGTTA AATGAGGCAC TTIGTTT

SEO ID NO:650: (Length of Sequence = 286 Nucleotides)

GGGTTGAAAG GAAAGGTGAC AGGAAAGATG TGTTTAGCAT CCATGAGCAG CTGGGGAGAG TCTTTCCTGT CTCGTAAACG CATCTGAGAA GATTAGGAAA AAAAATAAAC AGAGCATCAG TTCTTTGAAT CTAAAAGACT TINITCTACT AAAATTTCTA CCCTCAAATT CTCAACTAAT GAAGANTGTT TACTTTTGTT TTAAACTCAC TTCATTTTCC CAATTAACTA TTATCAAAAA AGTTAGTGCA TTGTAAAATA AGNTAATAAA GGNTAACACA TTATCC

SEO ID NO:651: (Length of Sequence = 360 Nucleotides)

GATAATGIAA ATTITIGCIT CIGGGCITGI CATCAGGATI GCAATTIINA GATTIAGIIT GCTAATIGIT TGGCCITTGA
AAAATTATAT ACACTIGGIT TGITTIGGIT TICTIAAGIC AAAACAAGGA AATAAAATCA CATTIGCITT CCAAGAAAAG
ATAATGITIA AGIGGITGIT TAGIGITTIG TGICITIGGG GGIGGGAGGG GGIGIGGGA ATACACAAAC ACACACACA
AAACACACAC AGICTATATA TAANCITATI GGAGCCATCA CIATATITTA AGGAAAATCN AAATAATCTA TIGAAGCITT
AAAATTAGGA ATTITIGATT TAAGCTAAGG AGCCTATTIT

SEO ID NO:652: (Length of Sequence = 353 Nucleotides)

GIGGIGGENI CCIGIAATCC CAGCIACTIG GGAGGCIGAG GCAGGAGAAT CGCITGANCC CIGGAGGCAG AGGITGCAGI GAGCCGAGAT CGAACCACIG CACTCCAGCC TAGGIGACAA GAGCGAAACI TIGCCGGCAT TIACACICIC AAAAGAITIA ACGCAATIAC AATCAAAAAA CACTIGICAT ATATAACACI TITICACAIG GAAATAAATT GGIGGITTAA GGIPTACAAT TCCITTGAAT AAAATITCAG TIATTAGITA CAAAATGCTA AGACAGATIG AGGICTCAAA GAAAGANCIT TGAGGAAAAT TIATGGITIT AAAGGGACIT TCACCAAATA TGA

SEO ID NO:653: (Length of Sequence = 224 Nucleotides)

AAGACAGGGA NIACTITATI CAAAACCCAT CACAGAAATG GACAGCITGG GICTGIAACA AAGCATTCAT GITTIAGNGC ATAGGICAGI AATIGIATAT GAGAGCATAC ACTGCIACAT ACAAATTAAC TGNTCAGACC ACAACTITIC AATGITTAAA ACAGNATAAG CITCCCIGIA AAAGCAGCAC CITTIGIGAC GNITIAACTI TAGIATICCT CICC

SEO ID NO:654: (Length of Sequence = 353 Nucleotides)

GICAACTCTA TITICCATAT GAATTATTAG ATTIGGIGCT GICTIGIGAA GIAACTIGAT ACGATAGATG TGIAGIATGA
ATTITGICCA CATGGITGIG CCCITGGCAG AACIGCACGT ACCIGAAATG GITCCCIAAT TITITICIAG TATTACIATC
CAACACTICC TCICATAATC ACIAGIGIAT TGIATAATTG TIAAGIGICC TTTATTCATA TATTIAAATT AAAAGAATAC
TCIGGIAGGA TITIGAGGGC CAATAGIGIA TITICCACTGI TIGAGGIATT AGGAGGGCTA TITIACTGATA CCTGIAGIGC
CTTCCCATIC TGGITTATCA TGCACCTCTA AAT

SEO ID NO:655: (Length of Sequence = 365 Nucleotides)

GAAACINACT TCACATTICT CCAGGGAGG ATGCTTTGGA AAAACIGCTC AGTGAGATGA AGCACAGATC TGCTTTTNAT
CCCTTTTGTA CCTTTTTAAA GACATAAGGT ATGTTTTGAC ACTGGAGTAT ATATGAGGGT TGCTAACGTT TAGGTTGAAA
GAGCTGCTGT TGTCCACAGC TTATTTATTT NCCACCCATT TTTGTCTCCT GGTCTCATCC AGTTACATTT CCTGGGATAT
GTTTTTGGAG GTTGCTCAGA TCACGGCACT AGAGTCCCTT TGGGTTTCTC CTCCCTCCTC TGTCTATTTG GCCTCGCCCT
TGACAAACAT TCCCCACATT CACAACCAGG CCTTTGGCTA AATGT

SEO ID NO:656: (Length of Sequence = 372 Nucleotides)

GTCATGAGTC TGAGACCAGC CIGGCCATCA TGGCAAAACC CTATCTCIAC TAAAAATACA AAAGTIAGCI GGGTGTGGTG GCGTGCACCT GCATTCTCAG CGACTTGGGA TGCTGAGGCA GAAGAATCGC TTAAACCTGG GAGGCAGAGG TTGCAGTGAG COGAGATOGC TOCACTGCAC TOCAGTCTGG GTGACAGAGT GAGACCTTGT CTCCAAAATA AAAGAAATTT ACTGCAAAGG GATGTTGCAT TTCAGGTGAA TGTATGTAGC CTTTCAGAGG CCGGGCTATT TATTAGATGT ATTTTATAAC TGAGGGTTCT AGGTAAACAC AAGCCAAACA GATCCACCAG AAGCCTAGAG CTGTGGACTC TT

SEO ID NO:657: (Length of Sequence = 334 Mucleotides)

GGITGIGGAA AAAAAAACCI CCAGATAAGA TIGIGCCIGC TICATITICI TGIGAGGCIG CCCAGACAAA GGITACITIC CTGATIGGGG ATTCIATGIC ACCIGATICA GATACIGAGC TICGAAGICA GGCAGIGGIG GATCAGATTA CCAGACATCA CACCAAACCA TIGAAGGAAG AAAGAGGGCC TATIGATCAG CATCAAGAAA CTAAACAAAC AACCAAGGAC CAATCIGGAG AGTCIGATAC ACAGAACATG GITTCIGAAG AGCCCIGIGA ACTICCCTGT TGGAATCATT CAGACCCAGA AAGCATGAGC TIATICGACG GATA

SEQ ID NO:658: (Length of Sequence = 286 Nucleotides)

ACAAACCAAC TGCATTTCTT TCTGGATATT GTTGAACAAA AATAGCATTC AGTTTACCCN CTAGTGCTAA CAGAAGNENC TCAAGCTGTT CCCCCATCAT GGGNGCAGCC CTTAACAGAG GGCTGCACAA ATCTGCAGTG CTGCTCTGGG GAAGGCTNCA AAGCACTTTT TTCCCAAGAA GGGATGCTGT TCANGTCTGT TAGGGGAAGC ACACCGNCIN TGCCTGGGCA CAGATGAACT GCCCTTCAAG GCAATCATCA TCTTTTTTCT AATAGGGAAG GTTTGG

SEO ID NO:659: (Length of Sequence = 321 Nucleotides)

GGICTITATA TGITTCCGAG ACAGGACTGA AACTCCCTGC CITCAAGTCA TITTCCTAAG TAGCTGGGAC TATAGGCTGT
TTCTTTTTT AAAGGAAGGA TTTTATGITT ATCATGAAGG
GAGACACTAT TAAAAAAAGG CAAATCAGAA ATTTGGAGAA CITAAAGAAT TAAGACAGAA TTGTACCCTG
TAACCATAAA TATGTAGAAT TTCTACCATA TCAATAAGGT ACAGTTCT GTTGCTCCAC ATCCTCTTGC ACGGTTGGGT
A

SEQ ID NO:660: (Length of Sequence = 302 Nucleotides)

TITIGITAAGG ACATAATGIT TITIGACTGGG GATCATGITT GGCTGATGIA AATATTAATG CCAAAATAGG AGCTAGGATG
AAAGTAACAC TGTAATTAGT AGTAGAATTT ATTTCATATT AAAATGIGIC ATGACGTAAT TITITATGGCT TGGCTCAAGC
AACAATTTTC AGAGTGCACC CTCATTGATG CTACTCACAG AGACGTGGAT GTGCTGTTAC TGCTTTCTAA CTCTGCCTAC
TACGTGGCCT ATTATGATGA TGAAGTTGAT AAAGTAAACC AGTATCAACG NCTAAGTCTA GG

SEO ID NO:661: (Length of Sequence = 249 Nucleotides)

AAAAAAAAAA ACTCTCAAGG GTCTAACTTT ACCCATCATA AAATAATTTT GGTGCAAGGG TAGTGGCACA TTTTATTTAT
TTGGGATACC ATGCAGATGC AACCTAGCCC CATTCTTTAT GCAAAGTAGA TTATCCGTGC ATTTCTTCTG CATTCMTAGT
GAATCCTTAC TGGGGNCAAC TCATTCCATT TGGCAACAAT CTTTAATGGN CAGGCAATAT ATAACATTGC TGAAGTCTCT
TAGCACTAA

SEQ ID NO:662: (Length of Sequence = 340 Nucleotides)

TTTTTTTTE GCAGCCTTGT AAGGAGAACT TCACCATTTC CCAGCACATC CCTATGTGTG CGCCTATTTT AATGCACCTC
TCTGAAACAG AGACCTTTTT GTTCACAACC ATAACTAAAG CTGGAAAGTC AGTCTTCAGG CAAGGCGAGG GAGGAAACA
TCCCCATTAGA ATTTTTCAG GAAAGACTTA TGGNAAAAAA TATCTTTCTC CCACCTCCTT TTATCCCCAT GAGACACAGT
TTCCCCACTGT AATCAGGGTA ATATGCATTT NTAAGINCTG ATATGTGATA CATTTATGTG ATGGCAAAGA TAAGTCTGTC
TTGCATGCAG GGTACTAGAG

SEO ID NO:663: (Length of Sequence = 325 Nucleotides)

CACAACAATT CIATGAAATT AGCIGGGGGG ATACIGICCI TATTITICAC AGCIGAAGAA ACCAAAGCIT TGGGAAGITT
GIGACITCIC TGAGATCACA GCIGGIGATA GAAGGAGCIG GGACACGCGC TIGGGITGAC TGGCITCIGG TITITGGITCT
CIGGCITCIA GIGCIGGAAG AAGCCCICIC TITCCCTTCI CITICCTCAG TAGCATCIGA CICITITCAT AAGCAAACAG
CIGIATAAAC AAAGCCCCCA TITITGGICAA GCACAGGGIG AATGIGATAT TIGITCCCAC AACCITATIC TNCACTCAAC
AGCCG

SEO ID NO:664: (Length of Sequence = 300 Nucleotides)

TTGCTGAGAG AGATGATGTT TCATGGGTGA TGTCTCTGGA AGAGATTGGA TAGGACCCAA GCACAGAGCA AGAAATTGGC TTTAGGCAAG TCAGATTTGT CTTATACTAG TTAGGAGTAA AGAGAAATGG ATGATACAGA TGCAGCTATG TTCGTAGGAG GGAAGTGGAG GGAATTTCTG TGTGATGGCT TTAGTAATGT AGGCAGCAAG GTCAACTACT GACAGTGAGA GGAGAAATTC GGGGAGGCTG GTCACAGTTT GAAGTAATAG GTCATGGGGA GGCAGATGTT TGTGGGTGGA

SEO ID NO:665: (Length of Sequence = 327 Nucleotides)

CAAATAAGAA CCCAGAGAGA GGGAGAGATT CACAGACAAT AGCTTAAAAA GTCTAGAAAT TATAGACCGA TIGAGGTCAG
CAAGAAACAA ATTATTCAAT ATATCCCCTG AGGGCTAGAG CCAGACTTTC CCCTATGATT CCAAAATTAC TICGCAGTTT
CATTAGGGTG AAAGGCAGTG CAGTCTCATG AGTTCAGAAA GTAAAGGTTG TTCCTTAAAA TITAGATAGA CTTGACAACC
ACTTAGGATG GCATTTTGGC ATTCTGTCCC TGCTCATCAA AGAAGTTGCT CAAATTTGTG GGNTAGAGGA ATGAGGAGCA
AGAAGTA

SEO ID NO:666: (Length of Sequence = 319 Nucleotides)

ATTCCCAAGG AGAGGCIGAG ACAGAGAGGC TITGAGCIGI TCCICAGCCC CCIACCCTAA CICCCICCCT ACIGITGATC AGGCIGGTCT CIAACICIGG ACCICAGGIG ATATGIGIGC CICAGCCICC CAAAGIGCIG GGATTACAGG TGIGAGCCAC CATGCCTGC CIGGGITTTA TCTTAAGGIC TTTGIGITGC TGITCCATCT GCATGAATAC ATTINCTTCA TTTACTTACG TCTTAGCTTA AATGATACCT CCICTTCTT CCTACTGCCA TTATCTTCCC TTGICACTCC ATACTCAGAT TTCATTGCA

SEQ ID NO:667: (Length of Sequence = 288 Nucleotides)

GGTGGCAGGC TGCTTGCANT NCAACGCCAG GNGTTTCCTG ATGGGTCAGG GTGGGGAGGC TGCACACCAC ACAAGGTCAC CCTACTCTAC CTTCTACCCAC CCCTACCACA GCCCTGAGGCT CACCACTCCC CCAGGGCATG GGACTCTTGA TAATTCCAAG TCCATGAAAC CCCTACAATTA TTGCAGTGCG TATGANTCCT TCTATGAAAG TACTTCCCCT GAGTGTGCCA GCCCTCAGTT TGAAGGTCCC TTAAGTCCTC CCCCAATTAA CTATAATGGG GATATTTT

SEO ID NO:668: (Length of Sequence = 212 Nucleotides)

TONTTICINT TICTIATCIA TCINCITCAC CATGIGICIT 033333CCTGG AACATAGIAG ATGCTCAATA AATATTGATT GAATGAATGA ATGAATAAAT CINCITACAC CICTCATGCT TCAAACAGGG AAAGGCTAGA TIATITAGAA GICTIGICGG GGATAATAAT NAGCICAGIG GAAGCCCTCT AGITCTCACT CGAGTITTCIC CC

SEO ID NO:669: (Length of Sequence = 281 Nucleotides)

ATCTITICAA CCCIATCAAT AAGATGITAT GAAAGATTGG TICTCITGIT TACAAGIAGI ATAGAATCIT TITTGATCIT
TGACTCIGIG CIGCCIATCI CATCAATGIT GITGCIATTA ATATCIGICC TITAACACTG GATGITGGGA TCTIAGIAAT
GITGCIGATA ATAGGATTIT CAGCAAAACT TCCATATCCC TIGAAGATAT GGIAGITITAT ATTACTATAT CGATAACAGT
TTTGCCIGIG GAGATTTGAC TAGITTTAGG TGITTGGAAG C

SEQ ID NO:670: (Length of Sequence = 234 Mucleotides)

AATAAAGTIG GGATATIIGA TIGITTICIT TICIGATCIT TAIGCIGACT GCAGTATCAG ATACCATITC ATIGITTAAA
AATCTICCIT TITITITITIT TITITITITIG CATTIIGCIC TITIGICATT GITICAAAGT CAAGIIGATG GCCNCAAAAT
TCCAGAGGCT AAGCAATGCA GAAGIITICAT CIACIGGCAG CIAGIITITAT TICITAAAAA TACATTAAAT TAGG

SEO ID NO:671: (Length of Sequence = 252 Nucleotides)

CCTGAAATGT AAATTGTTT TAATATATT AAGAGCACAC AGAAGTCTTG ATTTATAAAA AAATAAATAT ATAACATGAC AAATTTACTG ATGATCCTGG GGCTCTGAGG TCAAACTCTT TAAATGATCA GTGAAAACAT AAAACATCCA TGATCTGTTA ACACACACAG GGGCATATTC CAGTTGTAAA AAACAANTTC CTTGAAGGCT CAGNACGTAC AAAANTCAGT MTTTMTGGCA GAAAGCACAT CC

SEO ID NO:672: (Length of Sequence = 366 Nucleotides)

SEO ID NO:673: (Length of Sequence = 349 Nucleotides)

CCTCCCATCT TGGCCTCCCA AAGTGTTAGG ATTACAGGCG TGAGCANCCA CACCCTGCCT GGTTGTGTAC TCTTTTAAAT
ACTAAGTTTT TAATGTTAAA TGCTGCTTTT AGATACACTG TAAAAATACA CCTATCAATG AGTTTTTTTA TTAAAAACAT
TGCAATTGTA CTAGNCTTTA AATACTAAGC AATAATTCAG GCTTCAATGT TGGTTTATAG TTTTCTCATT TCTTTCATTT
AATACCTCTG TAAAATGAAG CAGTTACTTC CATTTTCCTG AGGTGAGATA AGTGCCCTGC ACAAATGTTA TAGGNCCAGT
AAGTGAGGAC TGGAGCTCTG GATCCTAAT

SEO ID NO:674: (Length of Sequence = 256 Nucleotides)

GCACTITIGGG AGGCCGAGGC AGTIGGNICA CCTGAGGTIA GGAGTITGAG ACCAGCCTGG CCAACAGGGT GAAACCNGIN TICGTCTAAA AATACAAAAN TIAGCCGGGC GTGGINGTGC ATGCCTGTAG TCCCAGGTAC TCAGGNGGCT GAAGCCAGGAG AATCACTTGA ACCCGAGGTG GGGCAGGNGG AGGTTGCAGT AAGCCAAGAT CGCGCCATTG CACTCTAGCC TAGGTGACAG AGTGAGACTC CATCTC

SEO ID NO:675: (Length of Sequence = 292 Nucleotides)

GAAGTCATIT TAGACTCTCA ATTITAAATT AATTITGAAT CACTAATATT TICACAGTIT ATTAATATAT TTANITCCTA
TITAAATTIN AGATTATTIT TATTACCATG TACTGAATTT TITACATCCTG NIACCCTTTC CTTCTCCATG TCAGTATCAT
GTTCTCTAAT TATCTTGCCA AATTITGAAA CTACACACAA AAAGCATACT TGCATTATTT ATAATANANT NGCATTCAGT
GGCTTTTTAA AAAANTGTTT GATTCAAAAC TITAACATAC TGATAAGTAA GA

SEO ID NO:676: (Length of Sequence = 392 Nucleotides)

ATCAAAGATT GCAAACATT ATTTGATCC TGGACTACAG TGTGGGGGATC ATTGCTATGT TGGCTTGCCT TTNCTATCCA
AATCTGAACC CAAAGTGCAG CCTGGTGTAG CCATGCAGGA AGATATGTGG GATGCTGACT GGGATTTGCA TCAAAGCYTG
TTCAAGGGAT GGACAGGAAT AAAGGAAAAT NCAGGTCATA GATTGAGTGC TATATTTGAN GTAAATACAG ACCTTCAAAA

CIGGITCCIT CIGGICIGAA ATTAGAGTAG ACTCGITCIG AGTACTIGGC AAATGACTAT TIGATICTICI GATTCCCIGG NCICCATGCT CACCAGATGC ATAGCAGGGA TCICTCCIAG NCACTCACAT CCAATTITCA GG

SEO ID NO:683: (Length of Sequence = 329 Nucleotides)

GATTITAAAT AGITAAAACA TITTITITAAA TOCATAAGIA ATTOTTACTO TACTOATITA TACACACATA TACTOACATG
TACACAGACA TACOTACACA CACACITATA AATACATGIA TACACAGAAT ATAGIAAGGI CITTITATCOC TITTOAATGA
AATAAATATT GIATTOTATA TITAGAATAA ATAATGITGA AAAAGIGATT TIGGAGAAAG GITGAAATGA TIGAGICTTA
AGIGIGICAA TGIATAATCT ACCCCTTTCT AAACATCGIG TITTAAGIAG TCATCITACT TCAGAAATTA GAGGCTCAAT
GIGITTAGG

SEO ID NO:684: (Length of Sequence = 281 Nucleotides)

AACATGGCTG ANTIGAGATT ACACTGCCAT GATACATTEN CTGACAGCAC TTCACATTIT CCCTGAGTTG GOGACAGAAA
TCACACTGCC CAAATACATT ATCTGATGGC TCCTCATGTT TCCCAAAAGT TAGGAAAGGA GGTTCTATAT ACATACATGC
ACAAGTGCAT ACACACACA ACACATACAC ACACACACAG TGCTAGATGA GATGTTGANT GNCATAAGGA AATGAAAGTN
CCATCTCTCT NITNCCTACC CCCTGCATCT GTCCCTTNAT A

SEO ID NO:685: (Length of Sequence = 324 Nucleotides)

ATTITIAATA ATTITIAAACT AGCTACAAAA TGTCAATCAC TTCACAAACT GACAGAGGAG ACAGGAGGAA TTTAATATTA CATGCTATAA TGATATTTAT CTCACAGTTT ATATTTCATT CATTTATATT ATTITITITAA AAGGTTTCIT TATCAGCTAC TAAACATCTC AGCAATTTGG TGTGCATAGC TCTAGATTAA GCAACAAAGN ATTGTACTGA TAACAAACCA CAGGGGAAAT GGTGGTTAGT AAGAGTCAGC CTTATAAAAT TTACATCCAC ACTGTTTCA CAGCAAGNIT GCTCTCTCCA AAACGGTGGN CATC

SEO ID NO:686: (Length of Sequence = 380 Nucleotides)

CEACGAGGAG GAGGAGAAAA TTCCCCCAGA TTCGGGCAGG CCCGCACCCC ACATTCCGTC CTGTTTTGAG AGGAGGAGGG
AAGAGAAATA AACGTGGCAG CGCATAGAAG GCCAGCAGGG AGACTGCTTT CCAGACACCT CCGGCCCACA CAGCCGTTCA
CCCCCCGTTT TTTCAGTCCT GGAAAAGGAA TTCGGGTCTG TTTTCCTTTT GGGCTCTGTG CAACTNCAGC TACAGTGGAA
AAAAGCAAAC TGCTCTTGAT CCCAGGCCCT GCCTAAGCCT CAGCAGAACT TNTAAGCCTA AACTINAAGA GCCTCACCCG
GACGAGCAGG CATNCCTTAA CCTTAAAGCA ATCCAGTTTC ACGGCCTGT TCAGTGGAAT

SEO ID NO:687: (Length of Sequence = 305 Nucleotides)

GACACTICCC CICITITATG GAAGCATAGI AAGATTITIC CITIATGGCG ATCATGATGG AGAAGTATAT GCTACAGGAG GAGAGGITCA AATTGCAATG GAACCTCAGG CACTATATGA TGAAGTAAGA NCINIGCCAA TTGCAAAGCT GGATAGGACA GITGCTGAGA AAGCTGITAA AAAATATGIA GAAGATGAAA TGGCAAGGCT CCCTGATAGA TTGTCAGTAA CITGGCCTGA AGGAGATGAA TTATTGCCTA ATGAGATTAG GCCTGCTGGA ACCCCTATTG GTGCGTTAAG AATTG

SEO ID NO:688: (Length of Sequence = 390 Nucleotides)

GAAGTCATAA GGCCTAAATA TTAATCCAGT CTGTGACAAC GACAAGGTGA ATACAAGCCA GTCTCTACTT CTCTGGGCCT CTGTTTTCTG CACTTATAT AAAGATTGGG CAAGATGGTC TAACTTAAAT TTTATGATTC ACTAACTTGA TTTTGTATGG GGCAGATTTT NCTTCGATGA AATATTAACA AATACANCAC TCAAATAAAT CAGCNATGGG GTGCAGATGA GGACTACCGT TTCTACAGCA AAATATGGGT GAACTCAGTA AGTGTAGGGA CACAGAAGTT AATGCTGACC TCTTGCATAG CATGTATGGG ATATTAAATC ATTTCCTGCC TTCCATTTCA GGGGTGAGGG AGGAACAGCT GTTCCTGAAC TCTTTTAAGG

SEO ID NO:689: (Length of Sequence = 315 Nucleotides)

GATTTAAGIG TIAGCATTIC TAAACITGAG ACTCITAACAG TAAAAATAAA GIAATCIGAA ACCIGITICC ATGGGIAAAA CACICIGCCI GGIATTCIIG TACACAAAAT TIACIAAATA TGIGAATATC ATAAAATGAA AATATCACIC CCITCAATTI CITIGGCCIT CACAAATICA ATGIGACIAT GATCCITITC AATAATACIT TCAATGACAT TGIGCITCIT TAGAAAAAATC

ACTIAAGITG TAGCATACAA TAGTTAACAT TAGTCCTTTT ATTGCTATGG TATATGCTAA TTTTTTTAAA AGGGG

SEO ID NO:690: (Length of Sequence = 291 Nucleotides)

TTAAAATACT CCATATATT NAGAAGCAAT TGAAAATGCA TCCATGTATG TNATTTGAGC GITACTAGAA ATTTATTTAT
ACAAATCCAT ATTAATGTGC TAATAAGTGA CAAATATATA TATAGTCATG CACTGAATAA TGATGTTTTG GTCAACGATG
AACTGCACAT ACAATGGTGG CCCCATAAGA TTAAAATAGA NCCAAAATTT CCTATGGCCT AGTGATGCTG TAGCCATCAT
AATGTGGTAG TGCAACCCAT TACCTTTTCT ATGTTTAAAT ATACAAATAC T

SEO ID NO:691: (Length of Sequence = 451 Nucleotides)

TTGAGCATCC GGAATATGGA GAAGTAATTC AGCTACAGGG TGACCAACGC AAGAACATAT GCCAGTINCCT CGTAGAGATT
GGACTGGCTA AGGACGATCA GCTGAAGGTT CATGGGTTTT AAGTGCTTGT GGCTCACTGA AGCTTAAGTG AGGATTTCCT
TGCAATGAGT AGAATTTCCC TTCTCTCCCT TGTCACAGGT TTAAAAACCT CACAGCTTGT ATAATGTAAC CATTTGGGGT
CCGCTTTTAA CTTGGACTAG TGTAACTCCT TCATGCAATA AACTGAAAAG AGCCATGCTG TCTAGTCTTG AAGTCCCTCA
TTTAAACAGA GGTCAAGCAA TAGGCGCCCTG GCAGTGTCAA GCCTGAAACC AAGCAATACC GTCATGTTTC AGCCAAGCCC
AGAGNCCTAA GGTTTACAAA CAAACTATGG NCCGGAACCT CCTCAAGTTC T

SEO ID NO:692: (Length of Sequence = 363 Nucleotides)

GATTTINTGA TIATTGATAT TAGAAATGIT TAAAATTAAG ATATTAACAT TICATGAAGC TGAGTGGTGA GCACACCAGT
TITATATTCT CICIATATAA CITTGIGTAT ATTIGAAATG TITTCCCATA AAAAGTATIT AAGCAAGTIT AGGAAAGAAT
ATTGATAAAT GAAATCIAGA GACCATCAAA AGCCAATTIC ACCATCACAA AGTATAATTG TGITTCAAAT ATAATTGAAA
TITGIGTGACT GITGCATATT CICITTITTG TIGITGITAA TGAAAGCATC TIAAACAGTT GCCTTTCAAA GCTGTTATCT
TIGATANTAA CATACATTAA CCTAACATTG TGGACTTCTG TTA

SEO ID NO:693: (Length of Sequence = 269 Nucleotides)

TTAAGGGTCC CAAGACTGCT CTAACAACAA CACCCATTIC CATAAATATG GNICAATAAA CACTTATICA TICTITATAA
TTAGACTCTA TIGTTAGAAT TGITTTAGGT TTATAGAAAA ATTGAGCAGA TAGTACAGAA GATTGCCATA TACCCCTCAC
CCACAGAATT TCACAATTTA CCCTGCGATT AAAGTCTAAT GITAATATGA TATATTTAGT ACAAGTAGTG GGATTATATT
GATACATTAT TATTAATTAA AATCCNCAA

SEO ID NO:694: (Length of Sequence = 330 Nucleotides)

GECATAGICA CITCCAGACA TEGITECCIC TOCATGIGGA GIAGGICAAA GICICCGICC TCCCTGGCCA GGIGGAAGCT CCAGAGGGAC ATGITTCAGC TTAGIACAAG GIGGCTGACA CIACTCCTCT GIAGGAAGAG GCTGGCTGGA GGIGAGGGCG CCCCACTCAG CCTGTACCCCA TCAAGAAGIA TTCAGAAAGG ATGICTCTGG CATCCACAAG ACIACTGGGC GAACCACACT GCAAAAAATGA AAACTAGCGT ACACAATTTA AATTGGTCTT AAACAAGCAA ATAATCCAGC CATTGGTGAC TCTGGGAATC TAGAGTGCAA

SEO ID NO:695: (Length of Sequence = 344 Nucleotides)

CACTGIGACG GATGAGIGGA TATTICTITG TACCCIGAGC TCTITCATCC TACCTIGGIG GICAAATGIG AGAGCAAGIG CITTGGGGCT CAGAGGGCAT CACTCCAAGC ATCTIGGATG GAGICIGITG TGGIGAATGI NCTIGGIGGC ATCTIGATCA AGGACTITIGI CATCATTAGC CATCAAATGC TTGITGGICC TTCTCAACCC TGIAATGITG ATACTITAAAA AACIGGAAAC ATCCTGACAG AAACAGICGA GAAAGIGGIT GIGIGAGCTC TGGITATCGC ATTACAGITA AAGTIGGCAG ATAGGITCTG TATTCAGIGC CCCATCAAAA ACAG

SEO ID NO:696: (Length of Sequence = 324 Nucleotides)

CTTGAACGTG GCAGATAAGC ATTTTGATAT GCTGCTGGAT TCAGTTTGCC AGTATTTTAT TGAGCATTTC ACATCGATGT
TCATCAGGGA TATTGGCCTG AAATTTTGTT GTTGTTGTT TATCTCTGCT AGGTTTTGGT ATCAGGATGA TGCTGGCCTC
ATATAATGAC TTAGGGAGGA GTCCCTCTTT TNCTATTGTT TGGAATAGTT TCAGAAGGAA TGTTACCAGC TCTTCTTTGT
ACCTCTGGTA GAATTTGGCT GTGAATCCAA TAGACACAAT AAAAAAATGA TAAATGGGAT ATCACCACTG ACCTCAGAGG
AAAT

SEO ID NO:697: (Length of Sequence = 341 Nucleotides)

AATTAATCAA TCAGCCATTT TGGTGGCCGA AATTTATAAG GCAAGTAATA CTTTTAGTTT CTTTGATAGA CACCATGATC AGAAACATAG TCTCTTTCTT AAAGGGAAAA TAGGAAGTCT TCTGAGTCAT AACAGATGCA TGCATAAATT TCTCTGAGTC TTCATAAGAA ACACAAGCAA GATTTCACAG AGGCAGTGGA ATTTGAACTG AGTCTTGAGA AATAAGCAAT ATCTGAACAT GTAGAATGCA AAATAAAGGA TAAGCAAGTG CTAATGCCCA GAGGGGTAAT ACATATTAAA TANCCANTAA CCAATTGCTA CTTGTGTTTC TTACACTAGA A

SEQ ID NO:698: (Length of Sequence = 317 Nucleotides)

GCAAACCAGG AGAAGCAGAA GAGCAGGGTA AACCCTGGGT ATAATTTGTC TAGACCCCCA TGTCTCCTTT AGTCTGAGTT
CTGACATAAT TAACTGTCTA TGAGATGTAC TGGGCCTTTC CTCATTGCTT TTTGATGCCA CCTCACTAAT GTAAACAAAA
CATTCATTTT TTCATCCTAT TTTTTCTTAC AGCTGCTTAG CACAGTCCTT ATGAAAAAAAT GAAGCCTTGA AAATGGTATA
TCCTCTCGAC AAAGCTAAGC CTGACAAGTT GGCTGCATTA CCTAGGAATT AGAGAAGAGC AAGGGCAGAT GGTGGGG

SEO ID NO:699: (Length of Sequence = 385 Nucleotides)

ACCAGGAGAT GGAGGIGCIC TAGACTGIGA TGCIGGGAAA GGATTGIGGG CIAGAAAAAG GGCICCCTAG GGCCGGCATA
TGGGCCACTG GGIGGAAGAG GGGCTCTGAG ACCCICACCC TGGAGCAGGT CATCACCCAC ACCGAAGAAT GAAGCGIGAA
TTCGGICACG CITAAAATGT TGAATTGITG GCAAAAGCCCC AAGITAATGA AATAGCATGG AAAATGGATG TGATGAGATT
TTTGAATTGT AATTAGATTA ACATTGICAC TAGITATCAG TCTGATATAT CITATAAATC AAACGTTGGG TTGATTTATC
TTTTATCACT TCTAGGGNCT TACTCCTAAC AGTAACTCAC AAACCCAGCC CCAAATCAGA GGCTT

SEO ID NO:700: (Length of Sequence = 315 Nucleotides)

ATCAGTTGGA TITGCAGAGG ATTGGAAGGC AGCACCAGGC AGGCTCAGAC TCACTGCTGA CAGGAATGGC TITCTTTAGG
ATGAAAGAGT TGTTTTTGA GGACAGCATT GATGATGCCA AGTACTGTGG GCGGCTCTAT GGCTTAGGCA CAGGAGTGGC
CCANAAGCAG AATGAGGATG TGGACTCTNC CCANGAGAAG ATGAGCATCC TGGCGNTTAT CANCAACATG CAGCAGTGAT
GGCGCCCAGGC TCTTCAGGNT GGGCCTGATC CCNCAGTGGT GCTTACTNTG CTGACTGTGT ACTTATCTTC CCCAA

SEO ID NO:701: (Length of Sequence = 387 Nucleotides)

GGCAGGAGAA TOGOTTGGGC COGGGAGGCA GAGGTTGCAG TGAGCCAAGA TOGTGCCACT GCACTCCATC CTGGGCAACA GAGCLARAGT CTYTYTYTÄÄÄ ÄÄTÄÄÄÄHAT AAAAAAAAAA GGTAGGTCTT TTCATCATTG TGTTYTCTRG CATTAGGCAC TGTAACTICC ACCTACTAGT AACTGAAAAC ACGCATGTGG GAACATTGCA CAGATGGATA GATGCAGAGA TGAAAGAAGG AAAGCTAAAA TATTINCCAC GTGAAAACCA TGCATCCTGT TCAGAAACTA ATTCTGCCTT CACGCCTTCC AGGAGCATGG GAGGGGTGTC GTCCTGGNCC TTTTGTGGAT GAGGGGGACC ACATGGTATT TCTACTGAAA GAGTTTT

SEO ID NO:702: (Length of Sequence = 397 Nucleotides)

CATCAAAAAA AAAAGGAGCT AACTAGATGC TGTCAATAAG AGACTCACTT TAGATCTAGA GACACAGGTT CAATGTAAAG
GGATGGAAAA ACATATTCCC TGTGGAAATC CCAATGAGGG TGCTATGGTT TTGCATGTGG TTTGTCCCCA CCAAAACTCA
TGTTTAAATT TAATTGCCAA TGTAATGGTT CTGGGAGCCT GGGCCTTAAG AGATAATTAA GATGGATTAA TGTCTTTCCC
ATGAGACTGG GTTAGTCGAG ACTCTTGCAA AAGCATGTTG TCGTTAAGTG GGTCATCCTC CTTTGTCCTG TCTCTTTTAT
ATACACTTCT TTCCCCTTCT ACTTTTCCAC CCTATTATGG AAGCACCGTG AAGCCCTCAC CAGATGCCAC CACCATG

SEO ID NO:703: (Length of Sequence = 374 Nucleotides)

ATACAGGGTT AGACCAAAGA GGGAATTCAA TGAGGCCTGA TGGATTTATG GACCAGAACA ACAGAGGGGT CTTGAAGGAA GGAAGATATA GAAAAGGCAA GGTTGTGGTT AGAGAGGAAA TCCCAGAGTT TTAGCTCTGG GAGGTGTAAT AATTTTAAAAA GAATAAGTCC AGGCCTGGCC ATGACATGGG AAGCTGAATC TCTGCAATGT TTTTTCAAAT AGCATAATGG ATATCTTTGA CTCCTTACCCT GAAGCCAGAA AATATTAAAC TTGCATGTAT AATCATACAA ATGTATGCAT ACCTATTTTA ACATACATTT AACATATTTTA TACTTATGCT TTCATATATT CTACGTGAGG TACAATATAC TCCA

SEQ ID NO:704: (Length of Sequence = 422 Nucleotides)

GGCAATGACA TAGAGATGAT AAGAAACAAC ATGGTTTGGT AGAGGGAACA TTTGATTTAG ACTCTGCCCA TTTTTAGCTG
TATGACTTAC ATAAGTCATT TTGTGTCCAA GCCTCATTTT CTCCCATATG AAAAGTGAAG GGGTTGGATT AAATGACTAA
AATCCCCTTC CAGCCCTATG AGCCCAATGT ATTATGATCT CTGCTTTGTT TCCTTCTTAA GAGGCTTCCT ACTATAAAAT
GTGACCTATT TACATTTTAA GTTGAAGTAG CCCACAATAA TGAATAATCA NTTTAGATTT TCCTCATCTC CTTTGGGAGA
AATTAAATTC AAGCCTCTAT TCATTTGATG TTTTACAACA AGCTTCAAAG TTGGGCCATG GTTCATTCAC AGTTTTGATA
TTTTGAGGAC ACCAATAAAA AG

SEO ID NO:705: (Length of Sequence = 229 Nucleotides)

GCTGCGGNTC ATAACAGCTG GACTCACGCC GNTGACAGAG TCTTGATCAG TCCTCTGGGA ACTAGACGTC AGGCCTCACAC CACTGTCTGC GCTGATCTGG GNCTTTTTCT CCTCTGTCTC ACCAAGGTCA AAGACAGGTT TGATTACTTC AGGCCTCTGT TTTTCCAAAG NTTTTTGCTT TNNCACTTCC TGGTGCTTGT TCCACAATTC AATAGATGCT ATAAAATTT

SEO ID NO:706: (Length of Sequence = 255 Nucleotides)

GAGGACTGIN TACCTCAGTC CTCTCTCTAA ACTCCTCAGC CTCCCAACAG GGGCCTCCTC ACCTGGGTTC TGAGTGTGIA CCCCTTTTAG AGAGTGAGAT GCCACCCGGG CAGCACTCGT TAAAGCTGGC CAGCACGAGT GACTAAGGGG AGAGAGCATG ACATAGACCT GGGTGGCAAC GGGGACCTCT GGAAGCAGGT GGGAGTAACA NAGGAGAGGG CAGTNGAGGG TAGAGGAAAA GGACCTCCAG AGGTT

SEO ID NO:707: (Length of Sequence = 324 Nucleotides)

CORNEAGTOT GCCACTOCAC TOUACCITOS TGACAGAGTG AGACTCOGTO TOCAAAAACA AAAAAAACAA AGITGAACTA TAAACTGAAT TOCTOCCAAG GITAGITCAG COTATGCCCT GGAATGAACA AGGACAGCIT GGAGGITAGA AGCAAGATGG NGTCAGGCCA GATCTCTTC ACTGITAACA TITTCTCAGT TATAATTTTT GCAAATGTGG TITCAGTCCC TGCATCCATA ATACCTAGAA ATTITGATAA ATACTTGITA AACAACCAAA AATAAAACAT CCACAGCAAG GANTOGACTA TAAGGCGTTG

SEO ID NO:709: (Length of Sequence = 264 Mucleotides)

GGCCCGGTT GCATGAGGCA CTTTGTCAAA ATGAGCAGAT ACGTATGAGC ACTGAACTCT TGAGTGAATC AACCAGAACT

AAGACCCAGA TCCACGCACT CAGGAACTTG CTCTGAATTT CAGTTTGACA ACAGAGAAGT AGAATATTTC TAATTAGCTA

ATATATATATA ACATTTTTTA ATCATCCAAA ATTACAGGCA AATCACTTAA GGTCCCCAGC ACTTTACGVT GNAAGGTCAG

AGAGANCCCC ACAAAAAAGG TGTT

SEO ID NO:710: (Length of Sequence = 366 Mucleotides)

ATTITIATIA TATACATATC AGRACICACA ATACGRIGGI TATTITAAGAT GGCTGRITTAT AAGRATAAAG CAGRITGAGC

AACACGGATT GGCCATTATT GRACITCAGA TGAAAAAATCC TTACATGCGG AATCAATGTC TTTTAAAATT TCAGATAAAG

AATTINCATT TGAGGRGACA TACAATTGTA AGGCCTCATT TTTTGTCAAT TTTAAGACAC CATTATGGT AAGANGGATT

AATTITNCCA TAAAATTACA AACACCCTCC ATGCCTTGAC ATTCACATGG AAAGGGCAGC ATAACCATTT AATCATCCAA

ATGCATATCA GAGCAAACTC CTAGGGCCTT TAGGGGGAGG

SEO ID NO:711: (Length of Sequence = 216 Nucleotides)

GAAAAGCAGA AAAAAGTGGG GAAGATTTC TATCTTGAAC TTGTGAGCTG GAGAATTACC ATTAGTAGCC CACTAATAGG

TTATGGCCGA TGAGTCCCTT CATAACACAC TGAGAGCCAC TTTTGACACT CCCAGAAAAG GCAGGTTAAC AAAACCCCTT

GATGGAAGGT TAGACCCTCA TTGCCCAGTG TACCCAAGCC TCTTTGAACC TTGCCT

SEQ ID NO:712: (Length of Sequence = 276 Nucleotides)
ATTITITIC CATAGRACGI ATRACTICIT CATGIGITAC CIGCIACACI AGAATTATGA CCCCIAAGAG GGAAGAGACT
ATGICAGIAT CATTGATICI NATIAACACC ATTATITAGA ACCATGCTIG GCITAAAGIA GIAGCIGCIC AGIAAATATI
TATCIATGIG TGAATTITIA AGINCITCCI TIATATIGAN TIAAAATTAG TCICITGIGI GCAGCAGICI GGGITTGICI
TATGITGAAA TACTIATGIN GACITCIACA TACATT

SEQ ID NO:713: (Length of Sequence = 354 Nucleotides)

AAACTTACA ACCIGCACAT TIGHTATGCA TACTAAATGG TGIGHTAAAA TTAGGGITTC TITGCCTCTC TACACTACAC

TAATCIGCCT AAAGGTGGIT GITTCATAIT TATAATGCTA ATTATCATAC CTACCTACIT TAAATTTTAG GIAGAAAATT

ATCIGATITA AATACAAACA TATTTTTCTC ACATTGAGTA ATATGCATAA TGIAGITCCA AATGTATTC ATTACTATAG

TCACAATATC CAACTAAAAA TTACGCTATC TAGAATTGTA CCANCCAAAA TCTCGTATTG GCAGATCTTG ACAGGCTGGA

CCTGCAAGAA TCTGGCTTGG AATTTTAAAC CCAT

SEO ID NO:714: (Length of Sequence = 349 Nucleotices)

CAGTAATTCT CTTACATCCT TCCCAAAAAT CAGTGTCTAG GGACTAGTTG ATCTGGATGA GTTATACATG ATATTTGACT
TINCATAAGT AGTGGAAGGT TTCACTAAGT AAAGATCTGA GTTTCTTGGT ATCTGACGTT TGTATACAGA TGGTGTCCAT
TTGCTCAACC AGACAGGAGT TAACTTGTAT TAGAATTGTT TTTNCTAAAG TNATGTTACC TGAGAAATTA AGGACTGCAC
CTGGTTTAAT GTTGCTTCAC TTATCCCACC CTACAGAGAC CAGCAAGGTT CTGCCAGGCC TCGAGCATCC AAGCATGATT
TTCCTGTGAC AAAATCTAAA AATCCAACC

SEQ ID NO:715: (Length of Sequence = 302 Nucleotides)

ATATTIGAAA AGATCITCAC CAAAGATATA TOGATAGIAA GIAAATATAT GAAAGGITTI CACIGITAAT GATIAAAGGA
AATGCAATCI TIGIACATGAA TIGITATAAC AGCATCATIC ATAAGAGCCA AAAGGIAGAA ACAATCCAAA TIGITCATCAA:
CIGATGAATG ANTACACAAA ACATAGIATT ATCTATATAA TIGAATATTA CITIGGCCATA AAAAGAAATG AACTIGGGCCA
GGCGCAATGA CITIACGCCTG TAATCCCAGC ACTITIGGGAG GCTNAGGIGG GCGGACTGCT TI

SEO ID NO:716: (Length of Sequence = 314 Nucleotides)

GIATTITIAG TAGAGACGGG GITTCACCGT GITAGCCAGG ATGGICTIGA TCTCCCTACC TCGIGATCCG CCCACCTCGG CCTCCCAAAG TGCTGGGATT ACAGGCGIGA GCACCTGCGC CCCACCCCAT TITGGIGIGA TCTCAGCTCA CTGCAACCTA CCCCTCCCAA GITCAAGIGA TICTCCTACC TCAGCCINIT GAGTAGCTGG GATTACAGGG GICTGCCACC ACGNCTGGCT GATTITCCTA TITTNAGITG ACACTGCATT TCACCAGGNT GGCCAGGCTG GICTCGATCT CCCTGACAAG AGGG

SEO ID NO:717: (Length of Sequence = 279 Nucleotides)

ATAAAAATGC TACAGATTIT TGIATGITGA TITTITATCA TGCAATTICA CIGAATTIGI TITTCAGITA TAACAGITTIT CITTATGGAGT CITTGGITTIT TNCCAAATAC AAGATCATAT CATCIGCAAT CAAGGATAAT TIGACITCCT CCITTCCAAT TTAGATGICC ATTATTITTIC CICCITGICIG ATTGCTCTAG CTAGGATTIC CAGGATAAT TIGACITCCT TGTCATATCC CAGGATCATG GAGGAAAGG

SEO ID NO:718: (Length of Sequence = 161 Nucleotides)

AAGAAAAAA CATAAATAAT ATTAGAAATG GAAAAGTTAT AAATCAACTA CAGCAAGGNT TTAAAACTAT TATGAAACAA ACCAAGTAGA AAGTAGATCT GCCAAACAAA AAAGGAAAGA NACTGTTTCT TTCATAAATA ANTGACAATG GGGGAAAAAG A

. <u>SEO ID NO:719:</u> (Length of Sequence = 220 Nucleotides)

GACAGAATT TITTITTITTEA GACAGAATCI CGCICTGICA CCCAGGCIAG AGIGCAATGG CGCAATCICG
GCICACTICA ACCICTGCTG TCACAAATAA ACATCAGIAA GAGCCAGCAG TIGCICTAGG ATCICAGICA GCAAGCTIGG
GGGCIGICAG GAAACCAGCA GICACCIGIT TCICCCICTC CCAGCCCAGG GCIGACCCCT

SEQ ID NO:720: (Length of Sequence = 347 Nucleotides)

AGANATGANA GCTACATTAN CGANANAGGA ACTIAGGANT GAGGTCATTA NATATANCTA ACTACATTTT ANATACOGAT
ATCATATATT TCCTGATTAG TATCAGGTAN ATATCTAGAC TCCTATCCTG NATTCCGGTC TCAGATANAN AGGTCAGAGA
CANTTACANG GANGATGCTT CATATTATCA GGTCAGTATA TACCTANTTA TGTGCACTGG NGAGTANTTT ATTCTTCATT
ATCATTTGTA NACATTGTTT TTTCACATTT TTGTAGTTGT CCATANTGTA AGCTTGTGGG TTTGATTATT GTTTTCACAC
CTGGATCCAG CTGGTTTANA CCTATTT

SEO ID NO:721: (Length of Sequence = 313 Nucleotides)

AAAAGATTTG AACAGATAAT TCATCCAAAA AAAATATGGG TGGGAAAAAA AGCACATGAA AAGATGCTCA ATATCATTAG
ACATTAAGAA AATATAAATT AAAACCACAA TGCAATATCA CCICGTATCT ATTAGAATGT CTAATATTAG CAAGACTGGC
CATATAGAGT GTTGGTGAGG ATGTGAACAA CTGAAACTCA TACACAGTGC AGGTGGAAAT GTAAATGATA CAATTTTTTT
GGAAAAGAGT TGGCTGTTTC TTCAAAAGTT AAACATTACA TCTGCCATAT GNTCCAGACA TTCCACTCCT AAG

SEO ID NO:722: (Length of Sequence = 266 Nucleotides)

SEO ID NO:723: (Length of Sequence = 370 Nucleotides)

ATTATTCATG AAATAATCCA TGTAACATCA CTTAGCACTG AGAGTTAACA AAGGCAAATG TTACCTGAAT AGGAGGAAACC AGAGGAAGAA CAACGAGGTC TCTTTTATCT ATGCTAAGCT TTGTCTGAAT AGGAGGAAAA TGTGTGGCCT GTTGGTGAAT TTATTGCTTT GTGGTAGTAA TGGATTYCC TAAAGCTGTT TCCCTCTGAT CATTAATAAT CCCTGTACAG CAAAGGACTA TTGTCCTTTG GTATGAGTAA ATAACCCTGT TGGAAGCACC GCTTATCTTC AGACCACAGC GCATACTTCT TACTGGAAAA TATAATGCAG GTGCCAACAC CCAAAGGGCA TGACCAGGGG TTCCCCTTCC

SEO ID NO:724: (Length of Sequence = 478 Nucleotides)

GGACACAACT GAAGTGTGGA AGAAATGAAA GGGCGAAGGT GTGTTTTGAG AAGGCTCTGG AAGAAAAGCC CAACAACCCA
GAATTCTCCT CTGGACTGGC AATTGCGATG TACCATCTGG ATAATCACCC AGAGAACAG TTCTCTACTG ATGTTTTGAA
GCAGGCCATT GAGCTGAGTC CTGATAACCA ATACGTCAAG GTTCTCTTGG GCCTGAAACT GCAGAAGATG AATAAAGAAG
CTGAAGGAGA GCAGTTTGTT GAAGAAGCCT TGGAAAAGTC TCCTTGCCAA ACAGATGTCC TCCGCAGTGC AGCCAAATTT
TACAGAAGAA AAGGTGACCT AGACAAAGCT ATTGAACTGT TTCAACGGGG TGTTGGGAAT CCACACCAAA CCAATGGCTA
CCTCTATCAC CAGATTGGGG TGCTGCTACA AGGCAAAAGT AAGGCCAAAT GCAGANTACA GGGGGATCTG AAGCTAGT

SEO ID NO:725: (Length of Sequence = 356 Nucleotides)

GACAGAGGAG AATAAATGGA ATAACTTAGT TTTGTGAAAG ACTCACAGTA TCACTTGGT TCTGGACACG GTTCGAGACC
TGGCTGTGGC TTGCTGTGGC CTTGAGAGCC ATCCCACAGC AGCAATGCTG TTGGACCCTT TGGCTGGGAC CTTCAGGACC
CCCTGCAACA GCACTGTGIN CCTAACCTGC TGGCATGATG CCCCTTTNTT GACAGGGCTG CATACAAGGC CAGCGACAAG
TGGCAGGCAG TGACGCCAGC CTGGATTTGC TGAGGGCACA CGCCATGCTT CCTGCAGTGC CAGTGCTCTT CTNGGTCCAC
TTTGCAGCAA GGATAGATGT GGTTCTAGAT CCAAGA

SEO ID NO:726: (Length of Sequence = 387 Nucleotides)

GIGGIAGAGI AAATCCIATT ATATCGAGAT ATIGGICAGG CAAGAATITI NCITTIAAAA TAATITATIG TAAATGAACC ATAAAATITI NACCITIGIG CCATCITCIA GGCIATAAAA TAGICITATA AAGAATCAGA TIGITAAGAG TATATGAAAT GIGGATATGG ATGIGGAAGA TCCATAACGA GGATGATGAA AGCACATTAA GAAGCTITCT GATGGGTACA AAAAATAGAA TGAAGAAGAT CTAGTATTTG AGAGCACAAC AGGGTGACTA TAGICAACAA TAATITATTG TGCATTITCA CATAACTAAA AAGTATAATT GGGATTGTAA CAGAAAGGAT AACTGCTTTG AGGTGATGGG ATACCCCCATT TTACCCC

SEO ID NO:727: (Length of Sequence = 348 Nucleotides)

TACCTAGCTA GCCCTCAACC TCTTTGTTTT ATGAATGGAA AGGCTGGGAC CCAGACAGGG CAAGTGACTC ACCCAGTGTC
ACAGAGCTGT TAAATGGCAG AGCATGATTG AATCGGGCCA TGACTACTTT CCTACATGAC ATATTGAAAC CAGTTTGAGG
CCTCGGTTTC CTCTCTMGCA AAACAGAGAT ACTAATG

SEO ID NO:734: (Length of Sequence = 374 Nucleotides)

TOGIGAAAGA AGAGAAGGAA ACCITGGICT GCATGGCACT TGGIACITIT GTATIGCCIC CATGCCCTCC ACIGCAGCTC
CTGCCCTGCT CTGTGTGCAT CCCTCATGAG ACTCAAGACA GATAACCTCT CCTTGCCCTT TCATGTCCCA GCCCTGGNTC
TTGGACTCAA CCATCCATTG CATCCCCATG GAGGATTCTG CCAGTCCTCA GGACTCAGGA GCAACCCAAG GATGTCCCAG
GGTCACAGGA AGACTTGTTG AGGGGACCCA CAGGGGTGCC CACAAATTAT CAGTCCATGG AGAAAAGTAG AGAGGGAGGC
TCAAGGACCT CAGCACGTAA GGGACATTTT GAATTCTACA AGTCACGGTG GGAT

SEQ ID NO:735: (Length of Sequence = 348 Nucleotides)

SEO ID NO:736: (Length of Sequence = Nucleotides)

ACACTCCIGA CCICAGGCAA TCCICCCACC TCAGCCICCC AAGGIGCIGG GATTACAGGC ATGAGCCACT GCGCCCAGCC
TACACACACT CITAATAGAA GAAATGAATA ATCAAAAAAT ATTATTGITG GAAAAAATGI TIGAATCITA TITITAAAAAT
AATTAACGNI TICAATAGGC ATGITGAACC TITITICGGC TACIGITTIC AGCAATIGCA GITGAATGAG TACAAAATGC
ACCACAGAAT AGAGACIGCT ATCIACCCAA ATATTGCTGG TIGITGAATC CATGGTAGGG AATTINCATG TATTGTTACA
ACCAGCTATA AATACATCCC AAAATATGTG TAGAGCTAAA ATAGATG

367

SEO ID NO:737: (Length of Sequence = 243 Nucleotides)

THATCATTC AAACTTCATT TTATACAACG AGTGCATACA CCACTGGGGG AGINICTGAC TGATGCGTGG GAGGGCGGGC
GGGGATGTCT NCAGCTATGA GTAGGGAGGA GGCGGGGGAAG CCCTGGGTGC TTCCTCTCCT CGACTGACCG CTGTGTGTTC
GTCCCCAGAG GAAGAGCGGN NGGCAGTCAG CCCCGGGGG GATGGCACAN TGGAGAGACG GACCTGCAGA AGTGGTGGCC
AAG

SEO ID NO:738: (Length of Sequence = 358 Nucleotides)

CGAGTCAGAG CTGGACAGCG GCGATGCCAT CTTTACATGG CCAGACCGAG AGAAGGGCAA ACTCCTGCAT GGTCAGAATG
GCTCTGTACC CAACGGGCAG ACCCCTCTNA AGGCCAGGAG CCCGCGGGAG GAGATCCTGT AGCCACCTGG TCTGTCTCCT
CAGGGCAGGG CCCAGCACAC INCCCGGCCA GTCCTCCTAC CTCCCGAGTN TGCGGGCAGC TNCTGTCCCA GCATCTGCTG
GTCATTTCGC CCTGACAGTC CCAACCAGAA CCCCTNGGGA CTTGAATCCA GAGANGTCCT CCAGGNAACC CCTCAACGAA
GCTGTGAAAT GAAGAGGTTT CCTCTTTAAA ACTGGTTT

SEO ID NO:739: (Length of Sequence = 400 Nucleotides)

CATTICTOGC CAGGCACGGT GCCTCATGCC TGTAATCCCA GCACTTTGGG AGGCCGAGGC AGGCGGATCA CGACGTCAGG AGATGGTCTA GACCATCCTG GCTAACACAG TGAAACCCTG TCTCTACTAA AAATACAAAA AATTAGCTGG GCGTGGTGGC GOGTIAGIAT TICCITAAAT AACAGGITAC AATAGAAAGA TACTGCCTGG AAGITATCCT TITCATTTTG GITCATTTTC AGITTITGIT TATGATTTAC ATAGCTGTTT AATTCATTTG CTTATAGTAC AATCCTGCCA TAAAGTATTA AAGCACAAGA TACCTGTTAT TCCCTTCAAC ATCTGCATTT TITCAAGNIT TTATACTCTA TATCCACAGT ATGTCAGCAG TTCTTGACTG

SEO ID NO:740: (Length of Sequence = 374 Nucleotides)

ATCGICAGAT TCACCAAGGT TGAAATGAAA TAAAAAATGG TAAGGGCAGC CAGACAGAAA GGTCAGGTTA CCCACAAAGG
GAAGCCCATC AGACTAACAG CAGCTCTCTC GGCAGAAACC CTACAAGCCA GAAGAGAGTG GGAGCCAATA TTCAACATTC
TTAAAGAAAA GANTTTCAA CCCAGANTTT CATATTCAGC CAAACTAAGC TTCATAAGTG AAGGAGANAT AAAATCCTTT
ACAGNCAAGC AAATGCTGAG GGATTCTGTC ACTNCCAGAC CTGCCTTACA AGAGGTCCTG AAAGGANGCA CTAAACATGG
AAAGGGNATA ACTGGTACCA GNCACTGCAA AAACATACCA AAATTGTAAA GGGA

SEO ID NO:741: (Length of Sequence = 290 Nucleotides)

AATTATTICA TAATAATGTA ATAAACATIC ATGAACATAC CCTATCAAGC AAGAGCTAGA ACCTTGGCAA TCATTTCCTT
GACTCCTCCA GITTGTGGCT ATCATGATAT TCAGCCCCAA GITCATCATT TCTGTITTTIN CITCIATACA GGITTCTTAT
ATGTATTTCT AAAAATCATT GGITATTTCA TCTTTGTAAA AAGTCATTGT NCTATTTTCC CCACTAGTTC TACATTGCAT
TCATATTGTT GTGGGTTGTG GTAATTCATT NATTTTGACT GCTGTATAAT

SEO ID NO:742: (Length of Sequence = 274 Nucleotides)

TTAAGAGGAA AAGTATCTTT AGGAATTINT TICTATAGAG TICTICATTA ACATTIATAC GAGTITTTIG CIGAGICAGA
TGGACAGITG GGTTCIGATG CITTINCCTT CCCGCCTGCC AGGCIGGCCC AGGCAGIGCT CCCACCANIC TATGAGCGIN
TCCGGGGCCCG NGGATCIGGG CAGCATCCAT GGTGCCGGGG CCATCCCCAG CGGNACCACA AGGINGCAGC GITGNICCAC
GAAANACCGN CITICCGCTC TGCTTCCCCA AAGG

SEO ID NO:743: (Length of Sequence = 398 Nucleotides)

TTGCTTTGCA GITATCTGGA ACTOCTOGIG CTCTTTCAGG AGCTCCTGGG TGTGCTGTAT ACTGGAGCCC GTGGAGGTGT
GTGTGGAAAG GTAGAACTCG CCATTGTCAT GGATCCATTC CAAAGCCTGC TTGGCACTCC TCTCAAAGAC CACGTACTGC
TGACACTGGT CCAGCCGTCT CTTCCTCATG GTCCAGTAAT GCAATACCCT GTTCTCCCGT TGGAAGAGTT CATTCAAGAT
ATTTTTCACT TGCTGTTCAG GAGCTTTGAT GTGCGTCACC ATTCCTGGCA TGTTCACGCT TGTTCCTGTG CACGTATTTC
AGGAAGACGT CTGCATTNCT CCGAGCAAGN GGTGCAAGCC TTCAGGAATG CCTCCTTTNC TNCAGGGTGC GGTTTTCA

SEO ID NO:744: (Length of Sequence = 359 Nucleotides)

TGGGACAGAG TCTTGCACTG TCACCTGGGC TGGAGTGCAG TGGTGCAATC TCAGCTCACT GCAACCTCTG CCTTCCGGGT
TCAAGCCATT CTCCTGCCTC AGCCTCCCAG GTAGCTGGGA TTACAGGCAC CTGCCACCAT GCCCAGCTAA CTTTTTGTAT
TGTTTTTTTT AGTAGAGATG GGGTTTCACT ATGTTGGCCA GGCTGGTCTC AAACTCCTGA CCTCGTGATC TGTGCGCCTN
GGCCCCCCAA AGTTCTGGGA GTACAGGGGT GAACCACCGN GNCCGGCTGG GGCTGCTTAT TTAAATCCCC TAGAAAGAGG
GATTCTNCAG CTACACCACA CCCTTAACTT NGAAGGACC

SEO ID NC:745: (Length of Sequence = 361 Nucleotides)

CCCTTAATTA AAAGTTTTAT TITTAAAAAA CGTAACAGAC CACTCTAAGA AACTTTOGCA TICAAAGCAG TAGTTACTGT TATTTGCTAA CICIGAAAAA AAAATTTINC CCCICACAAA CAACCGGCAA ACICCTGCCA CTTCCTAGCT TGGTGGCIGC CAGCGTGCAC TGCAGGGAAA CGGTGCGTGG AGGGATAGGA AGGCCCTCAC GCTCCCCAACC CACGGAGAAA NTGCAGATGG TGACAAGCTG CATCTGGACT CCAGGNTGTA TCTGACAAAG AGGGAGATGG TNTCCTCCNT CCCCTNCACC AGCTCCACTT TINCTGCTGA AGAAACAGAG ATGTGGAGGC AGGCGTGACC T

SEQ ID NO:746: (Length of Sequence = 285 Nucleotides)

GIGITITIAT TINIACCIAC AAAAAGAAAA CAAGATGAIG GIATCAAAAG GACAATITAC AAACTAAGAA TAGIAACATA
GCITICAGCA TCCIGIGCCI GAACATCACA CATCIACAAG TCITICAAGA CITAATGCAA CAGGAATNIG TCIGGAGACC
AGCAAGANCA TCAATAGAGA GCACTCAVICC CAAGCAAAAG CCACTAACCT TITAGATGAG AAGICCACAC AACGGATINI
TAGGGGAGGA TITIGGGAGAA GCAGCCCATT TGCTTAATAC ATTGG

SEO ID NO:747: (Length of Sequence = 302 Nucleotides)

CAATGCAGTT TTAGAGTGCT CATTCTTTCA ACTTATTTGA CAAATATTTA CTGAATGTCT GCCATAAGGC AGTAAAGGCA CAGAATGACT CAAAGCCTTT TINCCCTTAT GGGGTGTAAT TNCTAGTGGT GGAGACAGAC AATGAGCAAG TAAACAATCA ATCGGCTAAT GATAACTACT GTGAAGAAAA TAAAGCAGGN CAAGGGAATA GAGTATGCCA TCATTAAGAC TGGTTAGGGA AAGCTTCTTT GAAGACATGG CAGCTATTGA AAACCTGACT GATACAAAGA AGCAAGTCAT GT

SEO ID NO:748: (Length of Sequence = 346 Nucleotides)

GAGACCAGCC TGGGCAACAC ACTGAAACCC TCCTCTCTAA AAAGAAGAAA AAAATAAGAG TTTTGAGTIT TTCCAAAGAA GAATGCTCAG TACGTTTGIN ACCTATCAGA AAGAAGAATC TGGAGGTCCT GACGTGTAAA CAGAGTTGTG GGTACCATCT CACCAGAATT GCTGCCCTGA AGCCAAAGGA CTGAGCTGCT CAGATCTGGA AGTAATCTGA GCCCCCATTT CCAAGAAGAG AATTGCAGAA TTTTATAGGA AGAAGGGACC TGATCCCTGT CAATGGAAGC ATTTTAAAAT TTTTAACTGA AGTTCCAGGA GCATACAAAA AGCCAGGNAA TTTACC

SEO ID NO:749: (Length of Sequence = 325 Nucleotides)

CTAAACTITA TITICAAAAG CTTAAGGCCC AAATACAAAC TGAGGTCTTC CTTCCTAACA AATTAATACT AAAATGAAAC AGCTTTINIT GIGICCITAA GACAAAATAA GGAAGGAAAA CGTAGCTGCA GITGICCACG ATGGATATIG GITCITTAAA ATATATCTGA AAGTAGTAGT CAGAATGANI TATGGTTGGA AAACTGAGGN ATCTTCTGGT TGCAGGTGCA AAGTGACTTT MITTATTCTT GICTCAGTCT CCTTGATAGC CACTTCACTC TGCTACTACT CAACTTTCTC CTAAAAAATAC TTCATCTATT TTCAG

SEO ID NO:750: (Length of Sequence = 341 Nucleotides)

TGTATTTINA GIAGAGAAGG GGITTCGCCA AGITGGNCAG GCTGGICTCG AACTCCTGAT CTCAGGAGAT CGGCCTGCCT CGGCCTCCCA AAATGCTGGG ATTATAGGGG TGACACTGTC TCTGGITTAA GAGAACCATG GGCTGAGATA TINAGGAATT CTCCAGGCCA CGAATCTTGG GGCATGCAGC CTCTTCCGTA CCCCACAGCA TCTNGGGGAG CTGGTGTGCT GATGGGGTCA GCTCTCCCAG CTGCCTGGAA AATTCTCAGA CACTCCCTAA GAGGACATCT CCACCCCTNC CACTCTNACG TCACTGCTTT CTAACATTGC TCATTTGTTT G

SEQ ID NO:751: (Length of Sequence = 377 Nucleotides)

TTTTTTGAGA CGGAGCITNG CTCTGTCACC CAGGCTGGAG TGCAGTAAGC CCATCTCTGC TCACTGCAAG CTTACACCAT
TCTCCTGCCT CAGCTTCCCA AGTAGCTGGG ACCACAGATG CCCGCCACCA TGCCCGGCTA ATTTTTTGTG TGTGTGTTTT
TAGTAGAGAT GGGGTTTCAC CATGTTAGCC AGGATGGTCT GGCCCTCCAG CTTCCTCTGA GTCCCTTCAT AAACATTTGT

TTATCTTGTA AAATAATTTG TTCCATTTCT AATTAGTACA TAATGAGAGA GGCAGTGTGA TGGTTTGTGC CTAAGNCCTT
TCTTGCCAAG ACTTTCAAAG CCAAAAACTT CANCAGTTTT CCTAGATGAC TAGACAG

SEQ ID NO:752: (Length of Sequence = 359 Nucleotides)

AAGTCAGGCG TTCCTGGGGC AGCTGTCCTG TGAAGTTGGT GGGACGTGCT ACCCTGGGCC AGCTCCAGGT GAGCNTGGCT
TCGGTGGTCC CCGTGGGCTC CTNAGTGGCG AGGGTGAGGC CTGGCACTGG GCCTCTAACT GGCCCCGTGG CCCTGCAGTC
TTTNGTGCTG GTGTCCCGCT TGCCCTTTNT CCGGCTGTTC CAAGCGCTGC TAAGCCTCAT CGNCCCCNAG TACTTTNACA
ANCTGGCGCC CTGNCTGGAA GCAGGTGAGT GGCCATCANT CGTGGTCATC TTGGNCTCAT NATCCCAGCT TTGGCCCCTG
GTTGGGCTCG GCAAGCAGCT TCTCCTTGGG GAGGGTCCT

SEO ID NO:753: (Length of Sequence = Nucleotides)

AGCTTCAACT TGGAAAGAAG GATGATGCAG TITTGGGCCC TCCGGCCATC AATNACCGAC AGCNCTTTGA CCTTGCGGGA AGCCAGGTAT ATGINITCAG TGGAGCCCAG CTCTTTCTGG TGCCTCTGGT AGGCTGAAAA CATCTTTTCA AAATCCTCTA GGTCCAGGNT CCGAAATACC TGCATGTCAT CAATCTCATT CCATACGGTG CCAGGGACAC GCTCCTCATT CAGCTTCACC CAGTTGAAGG ACTTCAGTGG GTGAGAAGGC TGGGGGACAC GCTTTTTCCT GAGTGGGACG

SEO ID NO:754: (Length of Sequence = 342 Nucleotides)

CIGITGAAGI GCAGGITIGA TCCAGCCAGI ATAGAACIAG CICIGIAGGG GIGAGGAGGA CIGINCIGIG TATCATCCIT
GATTGINITC CITCAAGGAG CATIGCACIG TAAGIACATC AGAAIGACAA ATTGATGAAC TGCAACAGIA TCTITITIGIC
AATGITCCAC ATAATGCAAA TGCCATACGT TGTGIGAATA TTATGITGGA ATACAGIGCT GATATCTTGG AAAACCATAA
CTGCCICTIA ATITAACATA GNGIAATACA TAGINCIGIA TTTTTTTTTAA AGIGAGCINT AATGGGNAAG TATTTTTNAT
ATGCTTTAGC TATAGCTAAA GG

SEO ID NO: 755: (Length of Sequence = 321 Nucleotides)

CATTGCCATC TTCTCAGTCC TTCTCCCTTT CTTTCCAAGT AGTTTACGGC CCTAGGGCGA AGGTGGCTTT TATTTCCTCT CTTGGGGAAG GAGGGGGAG GAGCTTTCCC AAGCACATCA ACCTAAGGAA GGGGTGGTTG CNCCCCCAGC AGGGAGGGGC TGGAACTGCT GATCATTCGG AAGGAAGGGT TCGTTCTTGT CCACTTCCTG GCCCTTGGCT GCAAGGGTGT GCTTNGCAGG GGTCACTCCC CTTGGGGGTG GCAGCTCCTG CATCAGTNGA GGGCACAAGG AGGTATCTGC TGGTGTTCAC GAAGAGGAAGG

SEO ID NO:756: (Length of Sequence = 368 Nucleotides)

TGCATGIT GCATGINCT GIAATCICAG CIACITGGAG GCTGAGGCAG GAGAATTGCT TGAACCIGGG AGGIGGAGIT
TGCAGIGAGC CAAGATCGCA CCACIGCACT CIAGCCIGGG TGACCGAGCA AGATTCATTT TCAAAATAAA TAAATAAATA
AATGAGAAAA AAATATAGAT ATAGIAAAGG GAACAATTAC ATTCIACAAT ATTTIAGCAG AAGIAAATAT GGITTAATTC
AATGGAAACA GCICIGCICT AINGAAAATT CACAAATATT AAAAATAAAC ACACICIACA TIAAACCICT GAGCACTAGA
NGCITACCIA CITATICATA GGGCICACAT ACIGIAAGGG GGGIAAAT

SEO ID NO:757: (Length of Sequence = 339 Nucleotides)

CITCCACTGC CAGGITATCG TCCCGGGAAG CCCCCCACCC CCTCGNTTTC CTCCTCCGCT TTCCCTAACC CGTCTCGCGG
GGGCATCTAC GNCTCGTCCT CGNCCTCCTC CTNCTCGAAC TCCCCTTGTT CGTCGGCCGT GGCGTCCTGG TACTGCTGGT
ACTCGGACAC CAGGTCGTTC ATGITGCTCT CGGCCTCGGF GAACTCCATC TCGTCCATGC CCTCNNCCGT NTACCAGTGC
AGGAAGGCCT TTCGNCGGAA CATGGCCGTG AACTGCTCGG AGATGCGCTT NAACAGNTCC TGGGATGGCC GTGCTGTTTC
CGATGAAGGT GGCCGACAT

ACAAAGITIGG AAGIACICCI AATGEAGGGA AAGAITIGCCI AAAAGCCATA ATGAAAAGGG TAAATCATAA GGITCCACAT GITGCTCTGC AAGCACIAAC TCTTCTTGGG GCITGTGTGG CAAACINTGG AAAGATATTT CATTTAGAAG TATGTTCCCG TGGATTTINC AACAGAAGIA CGTGCTGTGA TTAAAAATAA GGGCACATCC TAAAGIATGT G

SEQ ID NO:765: (Length of Sequence = 329 Mucleotides)

TTGTCTGCTT GATGCAGGAG CTGAGGAGCT GCACAGAAGG TTAAAAGAGC TGTAACACAA ACAGGGCTGC AACATGCCCC
TTGCTCCCCA CAGGGAGAGA AGAGCTCTGG CCCTCGGAGA AGCCCAGACC TGGGAGCTCC TTGAGCCCGG GCTGTGACTC
CCTCTTTGGG GCCCTGGTTG GCGTCACTGC ATTCGCCAGT GCCACTGTTG GAAGCTGCTT GINATGCGCC TAGTCCAGGG
GGAAGCTTT
GCAAGCTTT

SEQ ID NO:766: (Length of Sequence = 321 Nucleotides)

GCAGIGGCAG GIAGATITIA TIGGCCIGGG ACACACAGGG GATACCCICA CCCACGATGG GGIGGGGGGI GIGGIGITGA
AGATATAAIC INAIGGICAC TIGIGGIAGA AICGCGGGIT CIGGCIGINI IGGATGAAGG GGAGCCGAGG GCCAGGITGG
CIGGIAGCIG CAAACCCGAC TITCCIGCIG GCIGCATCIG CACAGGGAGC TGGGGGGAAG CAAGGAGTCC AGGGGCIGGA
TGCAGAGCTT GAGICGGAGA AGCCAGICIG CIGGITAGCA TGINCCATCI GCITTINCAA GGNCAGGGCA CCACCAGGCT
T

SEO ID NO:767: (Length of Sequence = 313 Nucleotides)

ACCGCCCCTC TAGTTCACTA TTCTGTCCCC GGTACCCAGG GCATCATAGA CACTCAACAA CCATTCGTTG AATATGCAAT
TGGATGAAAT GAATAAACGA CCAGAGGAAT AATCCAGACA GAGCAGCAGT GGCCAAGGGA AGGGAGGATT GATTTATGGG
AGAAAATTAG GGGAATGAAA TCCATAGAAA GGGTTTGCCT AAGINAGAGT GATGACINGA GCCAGAAGAC ACCCGGGGGA
GAGGAATINT TTCACATGGT AGGAAAAGGG GAGGAGGGAG AGAGGTGGGG TGGTGGAGIN CAGCCTCGAG GCT

SEO ID NO:768: (Length of Sequence = 372 Nucleotides)

TCICITCICI GCCIGITIAT ATTCIGCACG TCCITAGIAA CCCCIGIGGC CCACTICITA CITAGGICIC TCCIAACATG
TATCIATGAC ACATIGATCC CIAACAGCIA TGATICINCI TATACTITIN CAGIAATTIA AATTTIATCA TICIACIGCI
TGITCAATAC ATCICICIAT GIAAATCITG ACTCCATAAT GAGGITTITA ACTTCGAAGG GGITGGAAGI TATCIGCIGC
CTIGGIACCC CCCCGCCATT ACACAAGAGI ACATTITAAG CACATTACAC CTGAGTGATT GINGIAAAAC ACAGATGCAA
TCITICCACC ATCCTCTAGG AATTCITCTG TGGGCTTTCC ATTGGGTTAC CC

SEQ ID NO:769: (Length of Sequence = 321 Nucleotides)

GCAGCCAGAG CTCCAAGGCT CCCCGGGGGG ACGTGACCGC CGAGGAGGCA GCAGGCGCTT CCCCCGCGAA GGCCAACGGC
ATGGAGAATG GCCACGTGAA AAGCAATGGA GACTTATCCC CCAAGGGTGA AGGGGAGTCG CCCCCTGTGA ACGGAACAGA
TGAGGCAGCC GGGGCCACIN GCGATGCCAT CGAGCCAGCA CCCCCTAGCC AGGGTGCTGA GGCCAAGGGG GACGTCCCCC
CCAAGGAGAC CCCCCAAGAAG AAGAAGAAAT TNINTITCAA GAAGCCTTTC AAATTGAGCG GCCTGTCCTT CAAGAGAAAT
C

SEO ID NO:770: (Length of Sequence = 364 Nucleotides)

3

TTAAATCAGG AAATGTGATG CCTCCATCTA TGGTTTTTGA AAGTCATCAG CCAGAGCTAA GGTAATGAGG ATTCCCTCCT
TCATGTTCAT ATGTCTTTAC ACTGTGCACA ACTGTCCCTA AAAAAACAAA CCCCTGGCCA ATTTCTCCAG GCTTATCGTC
TCCCCGGTTT CAGTTACATT TCAGCTTAGC ATTTTCAAAA TAACAATTTG TTCTTGGCAG CCTGTCTATA TATTTAATTT

ACCICICITG TTATCCCCAC TITTCATGCT CTATGTCCCA TAGGCAATTT GACAAAGACT GCTTTGACAA AGGATTCCTA
GACTTCTATC TCTACCTCTC ATCTGACTTG GGCGGAGGAT TAGG

<u>SEO ID NO:771:</u> (Length of Sequence = 357 Nucleotides)

CAGCTCACTG CAACCTCCAC CTCACAGGTT CAAGTGATTC CTTGCCTCAN CTTCCCAAGT AGCTGGGACT ACCGGTGCAC ACCACCATGT CCAGCTAATT TTTGTATTTT TNATTAGAGA CAGGGTTTCA CTATATGTTG GCCAGGCTGG TCTCAAACTC CTGACCTCAA GTGATCCGCC CACCTCGGCG TCCCAAAATG CTGGGATTAC AGGTGTGAGC CACCATGCCC GGCCTAAATT ATAGCTATTT TAGAATGTTG AAAGTAGTAT TATGTGATTT CAGTTTGCCA TAAATTTTTC ATATGGTTAC TAATTATTTC TNTTTTTTGTG GATATATCT: CTGGAAATCT ATTGAGG

SEO ID NO:772: (Length of Sequence = 359 Nucleotides)

CCTCTCAGGA AAACACCTAG ACATTATGTA ATGTATTGA AGATTAATGT ACCCTTTAAC CAGCAGTTGT GTACCTAGGT
ACAAACTTTG CAAGCACACA CGCATGINTG TNCCAAAAAG CACATACAAA AACACTCCTA ACAGCATTAT TTGTAATAAT
AAAATATAAG AAATTACCTA AATATCCATC GACTGCCATT GGTAGTATGG TTATACAATG GAATTCTACA CAGCAATGAA
AAGGAGCTAG AGCTACATGC AACAACATGG ATACAACTCA CAAACGTAAG ACTTAGTGGG AAAANGCTAG ACACAAAGTT
AACACCTTCT ATATGTGGGT TCCAGTTATA TAAAACCCA

SEO ID NO:773: (Length of Sequence = 361 Nucleotides)

GAGCCIACGG CAGAAAAAGA AACATCITCC TATAAAAACT AGACAGAATA ATTCTCAGAA TCTGCTTTGC GATGTGTGCG
TTCAACCCAC AGAGTAAAAC TTTNCTTTTG ATAGAGCAGT TTTGAAACAC TCTTTTTGTA GTATTTNCAT GTGTATATTT
AGAGCGCCTT GAAGCCTACG CTAGAAATGG AAATATCTCC CCATAAAACC AAGACAGAAG CAATCTCAGA AACTAATGTG
TGATGGCTGC ATTCCACACA CACCGTGGAC CATTTCTCTT GATAGAGCAG TTTTGAAACA CTCTTTCTGT AGAATCTGCA
AGTGGGATAA TTGGGACCTC CTAGAGGGCC TTCGTTGGAA C

SEQ ID NO:774: (Length of Sequence = 387 Nucleotides)

GTTTCGCTCT TGTTGCCCAG GCTGGAGTGC AATGGCGCAA TCTCGACTCA CCACAACCTC CGCCTCCCAG GTTCAAGCAA
TTCTCCTGCC TCAGCCTCCC GAGTAGCTGG GATTACAGGC ATGCGCCACT ACCCCAGCTA ATTTTGTATT TTNAGTAGAG
ATGGGGTTTC TCCATGTTGG TCAGGCTGGT CTTGAACTCC TGACCTCAGG TGATCCGCCT GCCTCGGCCT CCCAAAGTGC
TGGGATTACA GGCATAAGCC ACTGCGCCCA GCCAGAAGAT GCATGATTTC TTAGGATCAT ATGCTGTTTG TAGCCATAAG
GTAAATCATG TCTCTTCCAA TCATGACTTT TGGGAACTCC CTGAATAATA AAAATGAGAG TTGAGAT

SEO ID NO:775: (Length of Sequence = 401 Nucleotides)

GAATTINICI TICIGCATCE TICIGICATA AAAAGGGGTA CIACTATAGA ATAGAATGCA GGCTTAGGAC CCCCGTAAGC
TCACTGITCA ACCCAGCCCA GCAAACTGGT CAGITATAAA TITINCIGCA GGICCCIGAA ACAACAACAA AAAACTGGAT
GAGGITTCCC TCCCATCIIG TITITATGICC TIGGGAGCII GACCTTATAA CCATACOGCG GIACTITINC TIGGICTCTG
CCATCCAGGG AACCAGAATT TGGGGGGTTA TGICATAGII AGCTCTAAAA ATTATCTTGA GCAGITAAAA GCCTTTGCAA
GCTTAAAAATT GACIGCTGTA GGNTCCTTCT GGGGAAGGAG CAATGGGAAA CCTTNCCAAA GCTTATAGCT CANCCAGCTG
A

SEQ ID NO:776: (Length of Sequence = 345 Nucleotides)

AACACTGGGT AAGCACTTTG TATGINCTGG GCACTCTGCT AGAGATAATG TGTCTGGAAT TGGTGGGTTC TTGGTCTCAC
TGACTTCAAG AATGAAGCCG TGGACCCTCG CAGTGAGTGT NACAGCTCTT AAGGTGGGC GTCTGGAGTC TGTCCCTTCT
NATGITCAGA TGTGTTCANA GTTTCTNCCT TCTGGTGGGT TCGTGGGTCT CGCTGGCTCA GCNGTGAAGC TGCAGACCTT
TNCGGTGAGT GTNACAGCTC TTAAGGCNGC GCGTCTGGAG TTGTTCGTNC CTCCCGGTGG GCTCGTGGTC TCGCTGGGCT
CAGGAGTGAA GCTGCAGATC TTCGC

SEO ID NO:777: (Length of Sequence = 229 Nucleotides)

ATTGGGGGAA COCAAGCCCA NIAATGCTAT GGCTGTTGCA GACTTGTAGA GGTACTGCCT TCATGGTCTT NGGTAAGATC TGGGGAGAATT CCCTGGATTA CCAGGCAGAA ACTCTNATTC TCTTGCCTTA CTTCCCCCCA AACAAATNAG TCTCTCTCTC TCTCTGTCCT GAGCTGCCTA GAGCTGAGGG AGGGGGTGAC ACAAGCACAG CTATGTCACC AGGAAGCCA

SEO ID NO:778: (Length of Sequence = 361 Nucleotides)

CAGAAACTCA GGAATAAAGC CATTAACITT CAAAGAATAT GITGIGITGI TCGATATTIT CCATTCCTAA TCCACATCCA
CGITGGICAA GTAGAGCTIC CTACTCAGAA GCACAGCAGI TGCCATGGIG TTCTCTTCCA TCTGAAAGCA GCAATTITCC
GCAGCGICCA TTTACAGAAT GTGCCATATT TACTCAGATT CTAATGTATA TTAAATATGC TTTGGAAACT TAACAAGAAA
CGTGCAAGCN CTCAGTAAAG AAAAGTTGTA GAAAACAAAA ACTGAACAGC AGGCTTCTAG TTTCTCCTCT CCCAAAATGG
CCTTAGTGGG ATTCAAAAAT GGGAAGTGTG AATAAAACTG C

SEO ID NO:779: (Length of Sequence = 392 Nucleotides)

CCTAAGATGC CIGGCACAAT CAAAGACCIT TGGIGGCTTC CAGCATTIAT AAGGCAGAGT CCAAACACAC ACTTAAGAAT GACTTACTCC TCIGGCGGAC CCCACCATTC CCTCACCCCG CTTTGGCTCT GTCCTCTCGT GGAGCTGCCC CTGCCCCTAA ACACTGCCTC CTCTCTACCA ACCCGGACCA TATTTCCCCT CCTCCCCTCA CCAGGTCCAG CAGTACCCAC CACGTTTGTG GACATCTCCC CAAGGAGCTC TCACGTATCA GAAGCAAGGA GTTAGCCTTC AGCCCCACCT CTTGTGCTTA GGTCTACAGT GAGTTTCCAG TGATGCTTCC TACCGACTGC TTGGGGGTGC ACAAGAGTNA GGCCAGCAAG ATNCCAGCGG AA

SEO ID NO:780: (Length of Sequence = 453 Nucleotides)

CTCTCTATTT TCTCTTTTCC TTTTGACCTA CCATAGGAGA CAGATTGCTC ATCTCCAAAT TTCTCTGCTG TCTGCGGANT
GCCTGGTTTT CAACCTTGGT TAGGGTTTGG CTTAGGAATA GCATAATATC CCTTTGTGAG AGGTTAAACA CTTGAGTTAA
ATTTTGGAGG CCAGGTGGG TGGCTCATGC CTGTAATCCC AGCACTTTGG GGGCCCAAGG TGGGCAGATC ACGAGGTCAG
GAGATCAAGA CCATCCTTGC CAATATGGTG AAAACCCGTC TTTACTAAGA ATACAATAAT TAGCTGGATG TGGTGGCACA
CGCCTGTGGG TCCCAGCTAC TTGGGAGGCT GAGGCGGGAG AATCCCTTGA GNCTGGGGAA GTGGAGGTTG CAGINAGGGT
GAGATCGGC CACTGCACIN CAGCCTGGGN TGAGGAGGCA AGACTTCCGT TTC

SEO ID NO:781: (Length of Sequence = 306 Nucleotides)

SEO ID NO:782: (Length of Sequence = 443 Nucleotides)

GICTOGGGCT CCTGACCTCA GGTGATCTGC CTGCCTCGGC CTCCCAAAGT GCTGGGACTA CAGGCATGAG CCACTGCACC TCGCCTAATT CTACATTTTN ATCTACAGCA GACCTTTTAT CATAAAAGAA TTTCTATAAA ACATTTCTCA AAAGAAAATA

TGTATTGACA TICTATTITC TITCICCICC AGATACTATI TITMGGATTI NAAACATACA CAATACTTAG GAGACTIGIT
TTACICAGAG TGGAAAATIT TNCCAGGGAC AAAGTCAACA CAAMGAAACA AACAACAAAA AATAGCCAGA AAGAGAACAG
TTAAGTGCAG CICGGTGAGI CCCGGCAGII CCTTCCCGGC ACTGGCTCGI CCCTGGGGTT CTCAAGGTTC CATGCGGCCA
CAGCGTCCGI CCACCTGTTC CACGMGAGCC ACATGCTGGA ATT

SEO ID NO:783: (Length of Sequence = 350 Nucleotides)

CATTCAGGCC GGGCACAGTG ACTCATGCTT GTAATCCCAG CATENTIENA GACATAGCAG TAGGGACTAT CGACAAAGAA ACACACAGAG GGAAAAAGAA TTCCACATTT GGGAGGCTGA CGCATGAGGT TCACCTGAGG TCAGAAGTTC AAGACAAGCC TGGGTAACAT GGTAAAACCC CGTCTCCACT AAAAATACAA AANTTAGCTG GGCATGGTGG CCTGGGGCTG CAGTCTCGAC TACTTGGGAG GCTGAGGCAT GAGAACCTCT TGAACCCGGG AGGTGGAGGT TGCAGTGAGC AGAGGTCATG CTACTCCAA GCCTGGGGCA ACAGAGGGAG ACCCTGTCTC

SEO ID NO:784: (Length of Sequence = 265 Nucleotides)

ATTACTGAAA AATGGAAGAA AATATTIGCA AATTACACAT GTGAAAAGCA GTTAATATCA AAAATATATA AGANACTCAA
AGGACTATAC AACAAAAAAC AAATAACCAT GAAAAATAAG CAAAAGATAT ATTATAANINA TTINCAAAGA AAGACATACA
TATAGCTIGG CAGATAGATG AATATGGCTC AAAGTCAATT ATCATCANGG AAAGGCAAAC CAAAACAACT CTAAGATATA
AACTCACTCC TGTTAAANIG TITAA

SEO ID NO: 785: (Length of Sequence = 363 Nucleotides)

GIAAACNTIG AGAAATCGGA TGGTIGCIGI GICTGIGTAG AAAGAAGTAG ACATGGGAGA CITTICATTI TGINCIGTAC
TAAGAAAAAT TCITCIGCCI TGGGATCCIG TIGATCTATG ACCITACCCC CAATCCIGIG CTCICTGAAA CAIGIGCIGT
GICCACTCAG GGITAAATGG AAAAAAAAA AGAAAAATGA AACCAGGAGI TGGCAATTAC TTITTITTIT TITTAAAGACA
GAGTCTIGCT CIGICACCCA GGCIGAAGTG CAGIGGIGAG ATCTIGGCIC ACTGCAACCT CCACCTCCCA AGCTCAAGTG
AATTCTCCAT GCCICAGNCT TTCAGAGINA CIGGGGATTA NAA

SEO ID NO:786: (Length of Sequence = 291 Nucleotides)

AACAACAATC AGCCACAATG TGCTTTTAAG GATTTAACTG ATAGTAAAGA TAAATGTGAG TNITAAGAAT GGGATTTTTA
GACTAGGCTG ACACAAGGGA TCTTCTTINA ATAAAGGNTCT TGAGCATTTG TNITTTTTGGA GCTCATCCTT AAGGGCTGGA
CAGGAAGAAT CCTGTGTTAT GTGTGCATGT TGAGCAATGC AAAAAACACT CTGCCAAATC CTNGATACCA CATGGTCTNG
AGAAATGCAT GAGTGATTTA ACGCACGGNT GGGTGTAGTC ATTATGTTCC T

SEO ID NO:787: (Length of Sequence = 256 Nucleotides)

TATTTCIGIA TAATTTINAT TATGACCATA AAAATAACAA TGIAGICAAT AACAATTTAA TIGIACATTI TAAAATAATT AAAGIATATA ATTACACIGN TIGIAATAAA AAGIATAAAT GITAGAGGIG ATGGATACCT TATTTACCCT AATGTAATTA CTACACATTIG TAGGCCIGAA TGAAAATATG CCATATAAGG CATAAATATA TACACATACT ATATACCCAC AAATACCAAT AATAAATTTC AATAAG

SEO ID NO:788: (Length of Sequence = 322 Nucleotides)

GGTCCAATGA AGCTTCAACT CGTTTTCAGC TCAAAGCAGA CGGCAAATCA GCAAAAAGCA AAAATAATGT ATCTTACTGC
ATTACAGACA AAAAAAAAAA AAAAAACAGA GTGAAACTAG ANCTATTTTC AATAGTAGTT TTCTGACAGC TATATAANCA
AATATAGANG ACATTATGGA ATTAGTGATG TGAACGAGAA CTTGTCCATG TATCCTGCCT GCCAGCAAAG GTAGAGATTG

CTGINATATT TGIAATGGIT TACIATGAAG GCTGTTCCAT AACCINCAAT ATCCACTGNT CTTGGGTGGI ATACCAAGGA

- SEO ID NO:789: (Length of Sequence = 357 Nucleotides)
- TCAATGIGGC ATTIGITTT NITAGAAAAC CCCTTAGTAA GCACITCICT AACCCAGAAT AGACACIGGG TATCCICCAA
 GAGICCCATA GCTTCATTT CATCITCCAC CCTCTTCTGA GAGGGGGAGG CAGGGGATAG GGGIGGIGIC AGGCAGICIC
 CAAAAATGCCC CTCCTAGACC CCTGAGAGAA TTCATGITGC CAGCAATAAA CCAACAGCAC CTCAGIGGGG CATCANAGGG
 L CCCTCTAGGC TCAAGGCTAT TGCCAAAGGG CATTCCTGTT TTATGAGCTT CACGATGGGA ACCAAGGNAG GCTCTCGCAA
 GACTTCCTAG GGGCTTGGIC CTTCAACTTA TGCCCTT

SEO ID NO:790: (Length of Sequence = 366 Nucleotides)

TGGCCAGGCT GGTCTTGAAC TCCTGACCTC ATGATACACC CGCCTTGGCC TCCCAAAGTG CTGGGAATAC AGGCGTGAGC ACTGCACCCA GCCTTGTGG ATCTTTTAAA GTACAGTTCC CATAGATTTA CATTAAGAAT AAAAAAGTCA TGACATCTTG CTTTTATATG GCAGTTTACT CAAGCTTTTT AAAGAAAGAG CATTCATCTT GCTTTTACGT GGTTTTAGAA TGTTGAAAAC CTTTTGATATAA ATCTGAGTAA TTTACTGCAT TTNCCATTAA TTCAGCTTAG TTAGACTGCT GGNTCCAGTG CTTTGTTTTG CTGTCACATA TACCCTAATA TGCTTTTTAA CATATGACCA AATTCC

SEQ ID NO:791: (Length of Sequence = 317 Nucleotides)

AACAACTCCA ACCATAATGG AGAAGGAAAT GGCCAGAGTG GCCACTCTGC AGGGGGCCCT GGTTTTACGA GCAGAACTGA
GCCTAGCAAA TCTCCTGGAA GTCTGCGCTA TAGTTACAAA GATAGTTTCG GGTCAGCCGT GCCACGAAAT GTCAGTGGCT
TTCCTCAGTA TCCTACAGGG CAAGAAAAGG GAGATTTCAC TGGCCATGGG GAACGAAAGG GTAGAAATGT AAAATTCCCA
AGCCTCCTGC AGGAAGTGCT TCAGGGNTAC CACCACCACC CINACAAGGN GATATTCTAG GGGGTACTCA AGAGCAT

SEQ ID NO:792: (Length of Sequence = 258 Nucleotides)

GATCAATATA TCCAGGAATT TGTGAAAAGA TCCTAAACIT TTCAAACATG TCACAGGTAG TACTTGAAGT ATGCTTGGTA
AAATGTACCG GTTAAAGCAG TATGTTTCTC AGATAGCCTG AGATTTTATT TAACAATTAT GTATCTAAGT CTACTAATAC
ATTTGAGCAA AAGAGTGTTG GGTMCATAAA TAAGAMGTCA GTATTTCACT TAGATTATTT CAGAAACTTG TAAGTMCCTG
TAAATAGCTA CTCTGAAA

SEO ID NO:793: (Length of Sequence = 282 Nucleotides)

GGAATGACAT GGTCATTCIN ACTIAAAAGA AACATTITAG GITCACACIT GCCAAGITAG GAAGAAAACC AACCTIAGAT
CCCTTCCCCC CCACCAATAC TCCTTTCCCC AAACACCGTC CCCACCCGNC TCIATGITTA ATTGAATTIT TATTTGTGAT
ATATAGAAAA CCTAACCCAT GGCTGINATG CTGAGTGTCA TTTGGCTTCA AGCTCGAACC AGGGNACAGC TTGGCCTGGA
ACCCTGAGAC AAGATGCTGG CCTCANAAGG TGGGGGCTCA CG

SEO ID NO:794: (Length of Sequence = 330 Nucleotides)

GITGAGGCTG CAGGGAGCCA TGITCACCCC ACTGCACTAC AGCCAGGGTG ACAACAAGAA CCTTTCTOGG CGTGAACCCA

GGGGGCGGAG TTGCAGTGAG CCAAGATCGT GCCACTGCAC TCCACCAGCC TGGGTGACAG AGCAAGACTC CGTCTCAAAA

AAAAGTTTAC TACTCGGCTT TAATTATTTC GTTTCGGTTT TGGGTGAAAT NATTTTATTA CTGACTGGTT CCTTAGTTGT

ACAGAAGCCT ATTATCTTTA GAGAGACTCT TCATGGTAAT TAACTCAGAT TCTTATTTTG CCTGGGTGAA AGGANGGCAA

GTGGATCTAA

SEO ID NO: 795: (Length of Sequence = 332 Nucleotides)

GGAAATAAAG GTGACATGAA CTAACTATTC AATCATGAAT GGTAGAAAAA AATGAAAATG TAACGAGATG GGATCCGGGT
CAAAGTCAGG GGAGGTATAG TTGAAGATAT TGAAGGAGTC ATTATGATAC CAAAGAAAAT GGAAAGANAT GGTATCCAGA
TAGGTTATCC TTGGAGAGTA TCCAGGGATG TCTCTTTNCC TAAGACCTTA GAGAAGGAAA GGATGGCTGA TAATATAGGG
AAAAGTTGAC ATGGAAGGAT TAAATAATTT TTTNAGAATT CACGTAAGGN ATGATAATCT GAATTTCCAG GGCTAGGCTC
AGAAGCAGAA AT

SEQ ID NO:796: (Length of Sequence = 305 Nucleotides)

CCCAAGGGA CAGCCIGANC TCCCIGCICA TAGTAGIGGC CAAATAATIT GGIGGACIGI GCCAACGCTA CICCIGGGIT TAATACCCAT CICTAGGCIT AAAGATGAGA GAACCIGGGA CIGITGAGCA TGITTAATAC TITCCITGAT TITTINCTIC CIGITTATGI GGGAAGITGA TITAAATGAC TGATAATGIG TATGAAAGCA CIGIAAAACA TAAGAGAAAA ACCAATTAGT GIATTGGCAA TCATGCAGIIT AACATTTGAA AGTGCAGIGI AAATTGIGAA GCATTATGIA AATCA

SEQ ID NO:797: (Length of Sequence = 337 Nucleotides)

GECTGCATTA TGACAAGAAG TCAAGCTTCA TGACAGTTAG TATGGGCTGG AGTCTGCAAA GTCTGAACTG TATTCTCATA
GAATGATTCC AGGTTTCAGG GTGTTCCACC TGCCAGAACC CAAAACTACA ACTATGGGG ACACAAGGGA AGTTTTAGAA
ATCTCCCTCT ACACGCATTT CTGGTTTTCT ATTATTCCTC CATGGCAGCT GACAGATCTG GAAGTGNAAA TAGGGGATTC
TCAAAATCAA AGCCANGAAG ACACCTTGTG TGACACCAAT GGAGTCTCAG AGGGTGGGAA TAGAAGTGAC TINGNCCCAG
GCATTTGCTG GGAACTT

SEO ID NO:798: (Length of Sequence = 341 Nucleotides)

GAACCCTGGA AGGICTAGGC TACAGTGAGC CATGITTGCA CCACTGCACC CCAGCCTGGG TGACAGAGTG AGACACTGTC TCCAAAAAATA ATAGTGATAA TAATAATAGT CATTTATTIT AAGTCTACAT GCTGAGATGC CAGAACAAGT AAAATTGGAT TATAGATTCA AGCAGTATGT AGGTATACTT TCATAAACTG AATACTGATG TAATTTTTGGA TGATTAAAAAA CAGNCTTTTA GTAGGTGTTC AAAAATCTGG NIAATTCCTT TCATGACATT CAAACATTTA GGTGGCCTGT CTTTGTTTTT TTAGGTTATA ACTTGCAAAC ATTCANTGT T

SEO ID NO:799: (Length of Sequence = 322 Nucleotides)

TTTTTEAGIA ATGAATICAT TIAATATAAA CITTAGIATA GCAGAATACT ACAGGITACC CACATITAAC CCTAAAAACA AACAAATGAC AGGCACTTCA GIGAAATAAC AAGCCCATGI TCAAATATAA AATGCTAAAA GIGAGAAAGA AATTATGAAA ATTATATACCT TTAATTTGCA GACATATAAA CACITITGGI ACAGTACAGA TGCATGATGC CAAAAAGTAA AATGNICCAG TTTAAGCTAA CACATTCCIT GITTATACAG NITATTTINC TATAGCTCIC ATATAANANA AATATINCCA GCTCACACAA TG

SEO ID NO:800: (Length of Sequence = 405 Nucleotides)

ATCAAGAGIT GIGIGGICIA COGACTGAGC CIGOCAGATA ACCCIGIAGI ACAATITITIN CAGCATAGIG GAAAAGAAAG CCATGGNICI GGGCAGGICA GGGITTGANC GCIAGIGCNI TGIATIAATG ATCATGATGA TAGCIAGIAG ACAGGGCITA CCAGATACTA GGIGCICICI TAACIGCIIT ACATATGINA GITAACICAT TTAATCITCA TGACATCACC CCIGAGATAT GGGIAATATI ATAATGCACA TITIATAGGI GATGAGAGG AAGCACTIGC ACAGATTACT CCAGCITAGT TCATAGCAGA GCTGGGACTT TTAAATCAAG GCACIAGATG GITCCAGAGC TTITGIACTAC TCITCCIGGG TCTTTCACAG TCIGAGCTGG TCCGGG

SEO ID NO:801: (Length of Sequence = 408 Nucleotides)

CIGCGITCCA TGIAGCGICT TCCACAGINC TCIGITATAA GATGGITTGI TACATIGCIG CAGATATITC TGCATGICIC
TIGAGITTCI CAAGACCAGG GITGIATTIT TCCATGICIG TCGATGAAAC AGIACATGAC AAAAGAAGGI ACTIAATACA
TGITTGATAA ATTAATTACT GITTGGIAAA TTAATTATIG AAGGAAGACC CAGACTGGIT CIGATAAATC ATTGATTACA
TTTTACAAAT TIGGATAAAT TAGGGGAGCC TTGAGAAGIT AGAGCTCTAG GGAAGGITCC AGGGAACGIT TGAAGGATGI
GAAATATGGI TITCAAAATT CATAGITTAT TGCAGGATTC TGCAAGTGA GGCCAAAGAT GAGGAAGANG
ATGGGCTT

SEQ ID NO:802: (Length of Sequence = 343 Nucleotides)

ATGAGACTTA CTCACTATCA TAAGAATAGC TIGGGAAAGA CCCACCCCCA TGATTCANCT GGGICCCACC CACAACACAT CAGAATTATG GGAGCTACAA TITAAGATGA GATTIGGCIG TGGACACAGC CAGACCATAT TAGACTCATA ATTIGNCTIC TGCACAGTAA GANCIGGGCT GGGATACCTC ATAGATCATA AACAAATCCG CACCCATGAA AAGATTTAGA GAGTCACACA GGAAAGTCAA CAGAAGNCAG AGAGATGTGG GTCCTGGNCT TGCATGTCAT TAAGTGGTGG GNTCCTTCAG CTTTCACATN TTCAGGCAGT GGGGTCAAGA AAC

SEO ID NO:803: (Length of Sequence = 182 Nucleotides)

GAATGGCCTT NTCTAACGGC ATGTATGACT TGCATGANCT CTCTAAAGCT GAACTGGCCT CACCTCANCC TGTCTTGCTG GCAAATGCGG CCTTCAGTGG GAAAGTAAAT GGCAGCTGCT GINATTACCT GGTCGNTGAA GAAAGACAGA TGGCAAAATT NATGCCTGTT GGGGATGACA GC

SEO ID NO:804: (Length of Sequence = 312 Nucleotides)

TITATTIACT GCCGITGIAA AINATCACAA AACATATICA TIGICAAGIG AATGCACAGG CITICAAAGG TGATIGIATT CIGCAAGGIG GGGAATAGCC AACIACCITC TAAGGIGAAT GINCAGCCIG CCATITCCAA CCCCAAAACT CCTCTAGATT CTCAACAGGG CAGCITCIGC TICATGCCIC TNITCGGAAA GGICAGCCCT GIGIAGAAGG CTTAATACCA ACATGCAGAT CCACCTGAGA ATCACTGGAA TGCTCTGGAA TGCTTCCGGA ACCCAGTCAG GCTINCGGAA AT

SEO ID NO:805: (Length of Sequence = 411 Nucleotides)

CATGCAAAAT TCAGAATATA AAAAANIGCA GGGCCTGGIT GCCCACATAC ATTCCTCAGG TTAAGGTGGA TTTAAAGATG CCCAACAGAA CCCAATGAAT CAGAAGCTAA AAGGGACACT TCAGTGATCA GCAGACGCAT TCTCTCACGT AACAAATGGA GGGAAAGTGA GCCAACATTA ACTAGCGAAG TCACAAGGCT AGATTAGGGG TGTACAGAAA TCTAATTCCT GGTGCTATTT GCAACTACAT ATATTTAAAA TACANGGAGA TAAATACCCA GAACACATTA AGCCTACTGA TTTAAACAGA NCATTTCAAG ACTGCTACACA AGAAAGGGAA GGGAAGGTTAAAT GGGAAGGAAG G

SEO ID NO:806: (Length of Sequence = 287 Nucleotides)

GCATTINAGT GCTGATACAG ATACAGIGAG TICCIGCCCT TICCICTCCT NIATATIGAA GOGATIATAA ATGAAGCTCT
TTAAACATTC TGAGAICINI AAGITGATTT CTACATGAAC TCCAAGIGGT GITAATGACA TITTICAGAAA AGATGCTITTA
CTTAGCTGAC AAGAAAAAGT ACTCTGTAAG CCTTTATTTG TATGTGATAA AACAGAGTTG ATAAAATAAT CTACTATTAA
CTTATCAATG CAGTCTTACA GAATCCACCT ANTTACAAAG TAGATAA

SEC_ID_NO:807: (Length of Sequence = 369 Nucleotides)

GCAGATATA ACCTITICTC AAACATCTCI AATIGICIGC ATACCCCACT AATATIGGCI ACATAATACA TITATTITTIG
TCATTIGGGA CTAAGIGCCI TACTIAGITI TGINCAGIGT ATTCATTAAT TGAAGAATA CITATTCAGG ATTTCTATTA
CITAGITTIG CICAATATAT TCACTAATIG AAGAAATATI TATNCAGGAC TTCCATTATA TGAGCACIGG CCTTIGIGGT
ACAAAGATAC AACATGAATC TGAAACTCAA TITAATCTAG AAAGATTTAT TAATATAANC TCATCAGAAA AGCAAGNCAT
CTACIGIGAT AGCTACAGIA TTGGITAGAA ATGGAAAGAG AGAGCAGAT

SEO ID NO:808: (Length of Sequence = 361 Nucleotides)

CAGGCTTIGT ACCAGCOGC ATACTCTCCA AAAGATGTCC CATCCTTTN CTITCCTTTG CATTCTTCTC TITCTTCAGC
ATGCATCCAG ATGGGTTTAT TTTCATCATC TACAGAACCA AACTCCCTTT CATGGGCACG AGGAGAATC TCTTTGTACA
GTGTTTCTGC TTGCTTGAAC TTTCCTTGTT TCAAATAGCA GGATGCCAGG TTATTTTNCG TCTTAGCCAC GTTGGGGTCA
TCAGGTCCCA GTTTTGTCTG GTAGATCTCG AGGGCTCTTT GATAATAATA TTCTACTTCT TCATACTTGC CCTGGGTTCT
GGCACAGTAA AGGCCAAGTT ATTTAACTGC TTGGCAACAT C

SEO ID NO:809: (Length of Sequence = 353 Nucleotides)

CTAATTTATC TICATGTCCA GIGAGCAGIG TIGCGITTIT CCTIGIAGCA TITGGAAATG ATTTACIGGA ATTACAAAAC CTATTITCCC TITAAATTIC AGCITIGGCI CIGGCTGCIT TITAGAATAA IGCAAGATAA AAATCACACC IGAGGGCIGA AAACGGAGAG GGAATGGAG ACTIGATATT TAAGCAGCIT GAATGGITTIT CCNITINCTIT ATTITITAAAG AAATGCACIT GCCTATGATA CIGICTCCC AGIGAAATGA TITACICCCC ATTACICTAT IGATACANIA TIGIGCATGC TAGIGTIGTA TITCIATACA GIAGCITGAA AATTGATTAA CCT

SEO ID NO:810: (Length of Sequence = 296 Nucleotides)

GAGGICAATG CITCCCAGGC TCGAGITGAT GCCCACAGGI GIATTGIACG AGCATTGAAA GATCCAAATG CATTTCITIT
TGACCACCIT CITACITIAA AACCAGICAA GIITITGGAA GGCGAGCITA TICATGATCT TITAACCAIT TITGTGAGIN
CITAAATTGGC ATCATATGIC AAGIITITATC AGAATAATAA AGACITCATI GATTCACTIG GCCIGITACA TGAACAGAAT
ATGACAAAAA TGAGACTACT TACTITNATG GGGAATGGCA GTAGAAAAATA AGGAAA

SEO ID NO:811: (Length of Sequence = 493 Mucleotides)

CCAGGAGCTT CTCCTCTCTT GCCAGGCCTA TGAGCAGAAA CCTCAAATAA ACCCTGGGCA GAGAAAACCA ACTTAATGAA
GAGGACGTTG CTGTTTCCAC TGGCTTCTAA TTTTGCAGAT GCAATGAGCA CTTACGGCTT TTGCAGTGGT TCAGGAAAAG
GCAAGAAGAA GCAGATTGTC ATGTTCCAAA GCCCTCTGAT GGCTGCATGG AGCCAGCGGT GCTGTGACTT TTTTTTAATAG
ETTCAGTACC TTTNATACGT ATGTCCTTAT TTACTCTTTA TCTATGCTCT CTTCCTCCCA TCAGGCTGGG AGCTCCCTGG
GGCAGGTCTG TTTCTCCCCT CCAGTCCGGA NTTCGCAGGA GCTGTGCCTC CCCCATCACA CTTGGAGGCT GTCTNAAGGC
AGGGGCTGTG GTCTCGCCA TTAGACTNGA AGCTCCCCAA GGTTAAAGGT CATATCCTCA AAAAAGCTTA GAATAGCTTA
GGAACCTAGG GGT

SEO ID NO:812: (Length of Sequence = 337 Nucleotides)

AAATTCACAT ACTIGIAAGI NATGCAAGCA AATTCICACA TAATTATTT TAAATGCTAG ATAGITGGTA TAATINCAAT CATTITAAAT ATGITAAGAC TIGITTIGIA CCCTAACATG AGGICTATNC TGAAGAATGI NCCATGIGCA CTIGAGAAGA ATGACTGGAG TGINCITTAT ATGIATGINA GGICCAATTA GCITATAGAA TIGCNCTAGI CCTCTATTTC CTTATTCANC TTTTTGITTGG TTGITGINCT ATCCATTATT AAAAGIGGGG TATTGAAGIC TCCTACTATT ATTGIGCTAT CATCCTCAGC AAACTAACAC AGGANCA

SEO ID NO:813: (Length of Sequence = 310 Nucleotides)

AGGIGGCCTC AGNICAGCCA AGCIGACCIT GGCACTIGGC TGGCTTCINT AAGGCANTAG AGTGCCCACA CATAAGCNCA CCACCINICC CCACCICCIC CCITCICICC CATGCCACCC CACTIGCTTC CAAGGGCTIG GITTCCAAAG TNACATCCAG GGIGIAAGAG GITGGGGAAA ACGICCIGCA AGNIGGCTICA GGGATCINAT TCCATCAGAT GGICTCATGA ATACTGTGGG

AGATTAAATC CATCTCAAAA TAGGCAACCA ATGCTATATT CTGAATNINA GGTCTCTGGA CTGAGTCCCCA

SEO ID NO:814: (Length of Sequence = 361 Nucleotides)

GATTIGAGCC ATCAGAATIC AGCITTIGIA GATAAAGAAT ATGAACIAAT TGACTATGGA TGGAATTATT GIATATAGTC
AGCITGCTGA ATTATTGGIT AAGCACIACT AACIATATCT TGGIAAACTA TGGIGCAACT GAGCCACCCC CTAAAAGCAA
AAGACATTTA GCAGTTCACC ATATTTTGCA ATTAACCAAA TGAGAGCCTA TGAGANTGAA ATGNITICAG GTGGAGTTTG
ACAATACAAT TCATCCNIAA TATATAGGGN NAAATATTTC CTCAAAAATA ACATCTATGT GGIAGGNCCT TAAAAACGAT
GGATGNAATG CATGCAAAAAT TCTCTGGTAC ACAGACACAT G

SEQ ID NO:815: (Length of Sequence = 301 Nucleotides)

GAATTINACT CITGITICCC AGGCIGGAGI GCAATGGCAC GATCITGGCI TACCGCAACC TCCGCCTGCI GGGITCCAGC
GATTCICCIG CCCCAGCCIC CIGAGIAGCI GGGACIACAG GCATGCGCCA CCACGGCCAG CCAATTITIG CATITINAGI
ACAGACGGGG TITCACCATG TIGGICAGGC TGGCCICGAA CICCCGACCI CAGAGGATCC GCCCACCTIG GCCINCCAAA
GTGCTGGGAC TACAGGTGTC AGCCACCACA ACCGCNCIAA TIAATACTIC TIGAAATTIC A

SEQ ID NO:816: (Length of Sequence = 310 Nucleotides)

ATCTITAACA TATTAAAATA GACATGAGAA AAATGTGTCA TITGATAAAA TGGGGGAAAT GIAATAAATG ATTACCAGAA ATATAAAATT AAGCCGTATA TGCNCTTAAG TAAATCGAAT CTAGGCATCC TTAAAATGTA AAAAAGCNIG CAACAAGAGT AAGGCNGCCCA GAATGATGTA AATTACAGGA ATGGGGTGTA ATGTAACCTC TAGAGGAGGT GATGTTTAGA AGAAGCAAAG NGAATGCAAT GANGAAGCAA ACTTGTTTTA GGCAAATNCT CCTGGGAGTG GGACCAGGCA GCCCCCTCTT

SEQ ID NO:817: (Length of Sequence = 225 Mucleotides)

SEO ID NO:818: (Length of Sequence = 225 Nucleotides)

TTAAAAAAAC CIGIAGITIC ATTACCITIT IGAATAAIGN CATACAAAAA ATGIATITGN TITTITGIGC IGIGAGAATI GAIGITIGIA GATTAATAAT CATTITGITI AGAATTACAA AATAGITTIT AAATATIGIC IGAGAAAAGC CAAAGITAAT GCAACCNAGI GGAAACIGIA AGACCNITIG AGIATIGITI GITTIATIGG ATGCAITIGG ATITT

SEQ ID NO:819: (Length of Sequence = 280 Nucleotides)

TIGACIAGCI TOCIACGICA TIAAAAATIC TI'IAAATAGI CIGICITAAT GGCIGCAAAT TITGIOGIAA GICIGGGCIA
AAATCIGATG AAATGITITA CCIGIGGITA AGIAATTIAG CAACTCGIAT CITTITAAAA TATTACAACT GGGNATICIA
GTACGICACA AACATTIGIN ATATCATTIA TITTIGIGCCA TIGICIGIGC TATGAAATAC AGIAGAATGA AAATTTACTI
CAAAGCATIC ATINICITCC CCCAGGGNAT GAIGGCAAAA

SEO ID NO:820: (Length of Sequence = 328 Nucleotides)

CCAGTTAATT TIGIAAAGIT TATAGAGATG GITTCAGTTA GACCTGTGCT GTCAATACAC TAGCAATTCA CATGCACATT
TAANITIAAA TCTAAGITTA AATTTAAATT AAGITAATAT TAAATAAGAT TIGAAATGCA ATTCTCAGTC CTACAAGCCA
TGCTTCAAGT GCTTCATATC CATGTGAGGT TAGTGGCTGC TATACTGGAT AGGGCAAAAA GAGAACATTA TIGTAATCAT
AGAAATTCTA TIGGTAAGIT TATGGGGTAG TACATGGACT AGAATGTAGT GAGGTAGTGA GCTGTGGATG CAGAGAAAGG
NCACTGGA

SEO ID NO:821: (Length of Sequence = 310 Nucleotides)

TCAGCATTGT TITCTGTATG INTIGAGATG ATTATTTGGT TITCCTTTTT ATTGTGTTAA TITGGTGAAT TGCATCANCT
TTAGTATCTT AAACCAACCT TGCCTCTCTA GGGTAAACCT TATGTGGTCA TAATATATAA NCCTTTAAAT ACATTATTGG
ATINCTTTTT TTAATATATTT GCTGAGGATT TTTCATGACT ATAATCATAA GAGATATTGG CATATGATTT CCTATACTTG
TAATGNCTTT GTTAGAAGGA GTTTATATTA GCNTTTATNC TGGCCTCATA AAATGGGTTG AGAAATGTCC

SEO ID NO:822: (Length of Sequence = 372 Nucleotides)

GCCAGATTGT NITCCTTGCG AGCCCTGAC CCCGGCTACT CTTCACCAGA CACGCCCGG CTTTGGCCA CAACACAGCC
GTCCCACCCC TGGTTCCTTC ACCTTAGCAG TAGCAGTAGC TCTGGGTGGA GTTGCCAGAG GAGCTGACAG GCCCTCTGCC
ACTGCTGCCA CCCCCAGGGC TAGGGAGGA ACAAAGAGCC TGCTTGCTGT GCTTGCACAT CCAGCATGCC ACAGCTGCAC
TACGGAGAGA AGGTCAGACA GTCCCCCCAA CAAGACCCCG ATCCCTCTAC TCTCCACCAG GGAGGGCCCT GGGCTTTGCG
CCCACAGAAC AAAACGTTCC ANCCCGGGCT GATCATTCTG GGTTGGCAGC GG

SEO ID NO:823: (Length of Sequence = 288 Mucleotides)

AGCTGGCATC CCTGGGGAAA ACCAACGAAC AGTCTCCTCA CAGCCAAATT CACCACAGTA CTCCAATCCG NAACCAAGTG
CCCGCATTAC AGCCCATCAT GAGCCCTGGG CTNCTTTCTC CCCAGCTTAG TCCACAACTT GTAAGGCAAC AAATAGCCAT
GGCCCATCTG ATAAACCAAC AGATTGCCGT TAGCCGGCTC CTGGCTCACC AGNATCCTCA AGNCATCAAC CAGCAGTTCC
TGAACCATCC ACCCATCCCC AGNGCAGTTA AGCCAGCCC AACCAACT

SEQ ID NO:824: (Length of Sequence = 325 Nucleotides)

CTCCTGAGGT CAAAGCTGCA CGTGGGGAAG AGAAAGACAA GGAGACCAAG AATGCTGCCA ATGCCTCTNC ATCCAAGTCG
GCCAAGACCG CCACTGCAGG ACCAGGAACT ACCAAGACGA CCAAGTCATC TGCTGTGCCC CCAGGCCTCC CTGTGTATTT
GGACCTGTGC TACATTCCTA ACCACAGCAA TAGTAAGANT GTTGATGTGG AATTTTTCAA GAGAGTGCGG TCTTCCTACT
ACGTGGTGAG TGGGAATNAC CCTGCTGCTG AGGAGCCCAN CCGGGCTGTC CTGGGACGCT TTNTTTGGAA AGGAAAAGGC
TCAGT

SEO ID NO:825: (Length of Sequence = 318 Nucleotides)

AATCAGCCCT ACAGCGATTC CTCCACCCCC ATTAGCAAAT ACCGTAATAT ATCACCTAG TAATCATCCT CTCACAATTC
TACTTTTCCT AATTINACCG TGAGTCAAGT TTCTTGACCA CAATGTTATG CTGAGGAAGA TCTAATGTTT TCCATGGAGC
AGAAATTGTT AGTCCTCAAC TCCAAGGTCT GCCTTGTCAA GCCCTGTTTA CCGTGTCTTC ATAAACCTTG TCAGGCATTT
ATTTATTCAG CACATATCTA CTGINCTCTG CACAAGAATT CATAAGGTTC TGATGAATTA TGTCCCTTCT GAGTGGGA

SEO ID NO:826: (Length of Sequence = 287 Nucleotides)

TACAGACTCA GGTTATAGGG TGINATTTYC TAAGICAATA TYCAGTTYCA CAGCCAGAAT CTGTGAAGAG AGAACAAACC ATGAGAAAAC TAACANTTYT ATGGTGATTG AGAGGTTCCA AGINCCTGGN GTTTTAAAAA AATCAGTTYT TAAAGATAAA

CAAACTAAAA CTAGTCCAAG CACTGAGACA GAGTATTAAA AGATGGTAGC ACACCCAAAG NGCACGGTGG GTCTTGAATA GCTAACATGT TTCAAGTAGT GGAGGNAGAT GTGCTTAAAT AGTTACC

SEQ ID NO:827: (Length of Sequence = 426 Nucleotides)

TITITITIGI TITIGGACAG AGICTCACTC IGICACCCAC GCIGGAGIGC AGIGGCGIGA TCICGGCICA CIGCAAGCNC
TGCCICCGG GITCATGCCA CICTCCIGCC TCAGCCICCA GAGIAGCIGG GACTACAGGG GCCCGCCACC ACGCCCGGCT
AATITITITIG TATITITIAGI AGCGACAGGG TITICACCGIG TCAGCCAGGA TGGICTCGAT CICCIGACCI CATGATCCAC
CTGCCICGGC CTCCCAAAGI GITGGACTAC AGGCATGAGC CACCGCGCCC GGCCGGATGG TTAAAACATT TTAAAAATAA
ATATITAGIG CTAAGACAGG ATATGGAGCA ACAGGAACTC CTATATGCTT GCTGGIGGGG AATGCAAAAT GGGIACAACC
ACTITITGGA CAAACAGITT TAGIAA

SEO ID NO:828: (Length of Sequence = 402 Nucleotides)

GECTGCTTGC TCCACTCAAA CAGGIATCTG GGAGCCAGCA CTCTGGCAGT CCTTCTAAGC TCTAACTCTG GTTTTACTGT
TTTNNAGGIG AAACCTTTGT CCTGGGGAAT AGTCTGGCCC GCTCCTTGGA ACCACACTCA GACTCAATGG ACTCTGCCTC
AAATCCCACC AACCTTGTCA GCACCTCCCA AAGGCACCGG CCCTTGCTTT CATCCTGTGG CCTCCCACCA AGCACTGCCT
CAGCTGTGCG CAGGCTATGC TCCAGGGGIA AGCTTACCAG AGTCCTGGCC CTNCTTCCCT CCCTCACTCT TTCCTTCACT
TCCTTCCTGA GCTCTGGGAG GCCAGAGAGG ACCTAGCTCT GTTGCCCTCT GNCTNGTGGT GGGGACTAGG GACTGGACTT
AA

SEO ID NO:829: (Length of Sequence = 417 Nucleotides)

ATCGGITAGG AGTCGCCITI AIGIGGGAAG AGAGAAAAAA ACTIGGIGAA AIGCITICIG GACTAATIGA AGAAAAATGI AAACTACTIG AAAAATTIAG CCITATICCA AAAGAGIATG AAGGCIATGA AGTACAGICA TCITIAGAGG AIGCCAGCIT TGAGAAAGGGG GCANAGAAGC ACGAAGIITIG GAGGCAACCI GIGAAAAGCI GAACAGGICC AAITCIGAAC TIGACGATGA AATCCICTGI CIAGAAATAG AGITAAANGA AGAGAAATCI AAACACTCIC AACAAGATGA ACTGATGGCA GATATITCAA AAAGGATACA AICTCIAGAA GATGAGICCA AANINCCCIC AAATCCACAA ATAAGCITGA AGNCCAAAAT CATTCINGCA AGGITTCITC CCAATGG

SEO ID NO:830: (Length of Sequence = 404 Nucleotides)

GETTIGAGAG TAGAACAGGA AGTIGIGAGI AGAGCCTIGA AGGAAAGAGA ACAGCAGGIG CATGENICCC CAGGCAGGAC

ȚCAAGGIAGC CACTCAGGCA TCAGAAAGAG TCAGGCGGCC ATGATGGCTC ACACCIGIAA TCCCAGCACT TIGGGAGICT

GAGICGGGIG GNICACCIGA GGICAGGAGI TCGAGACCAG CCIGACCAAC AGGGIGAAAT CCCTICTCTA CTAAACTACA

AAAATTAGCC AGGIGIGGIG GCACATGCCT GGGACAAATT TGGGATCAGI GITCTCCAGI CTGAACATAG TCTICTGITA

CCTGGGAGAG AGTGGICAGG TACTICCAGC TTCAGGGCAG CCAAAAGCAT TGACAAAACG ACAGGIAGGA TGGGGGGAGI

AAGT

SEO ID NO:831: (Length of Sequence = 330 Nucleotides)

AATTICACAG GITGIGICIT CIGAAATCIG TACCITCITA CICATAACAT TIAATGIAGC AITTICCAAC CIGACCAATC
TGCAGAAAAT ATATGICATA TATTAATIGI GIATACATGA ATATATGCAT TITCCTGGIA AAAAGICATA GITTINCATA
GATGICATGI AATCITITAA GAGATICICA AATAGGAACA TGATTCCACC CCAATAATGG TGAAAAATGA TCAATITAGA
TGAAAGGGAC CICAACAAGC CICTIGAGAT ATGAANCATA AAGAGNAAAT ATAAGCCGCA ACTITITGAC ATGACAGATI
CATAATGGIT

244

SEQ ID NO:832: (Length of Sequence = 402 Nucleotides)

CIGITITCIC CITITGITIT CCIATITAIN CICCCAGIGC TAACITGATA TCINCTIGIG TGIACACGIG TGINIGIGIG CAAATATAIT TCIAGGAACA AGAGCAAACA TICIAGTAAC TATCATICIC TGATGIGGAG AACITGGGCA GAGATCIGAG TIACAGCITT GIGGATTIAT TCICCIGGAT GAGAGGAACG CCCITAGAAT GICATGGTCC TAACCCCGTC ATGGATACCA GGGGTGAATG GCAGGGTTCI TCICCIGCCC AGGAGGAAGG GTATGGGGAG CCGGTGCATC TIGACTGTCA GGICACCTGT CITACCACCT TIACAGCTAG GCTTTCIGAG GIGCCAGCGI CICCIGGGAA TICAAACTGI AGITTAGAGG CAAGCTGGGT GA

SEO ID NO:833: (Length of Sequence = 398 Nucleotides)

AGCCTTTTIC CAGAGATCAG ACCTCTITAG ACATCIGAGA NITCATACAG GAGAAAAACC TIATGANIGC AGIGAATGIG
GAAAAGGCTI CICCCAGAAC TCAGACCICA GIATACATCA GAAAACTCAT ACCGGAGAGA AACACTATGA ATGCAATGAA
TGIGGGAAGG CITTCACAAG AAAATCAGCA CICAGGATGC ATCAGAGAAT CCACACGGGA GAGAAACCIT ATGTATGCNC
TGACTGIGGG AAGGCCTTCA TCCAGAAATC ACATTTCAAC ACACATCAGA GNNITCATAC TGGAGAAAAG CCGTATGANI
GCAGTGACTG TGGGGAAATC CITTCACTAN GGNAGTCACA ANCTTCCATG TGCATCAAG GNTINACANC CGGGGAGG

SEO ID NO:834: (Length of Sequence = 394 Nucleotides)

CITITITIGIT AGICIGIAAA ATCATITCCA GGIAAAATCI AGAGCITAAT CCATATGING IGCCATCITI IGCITITCCA CACCICINAT CCIAGGIAAG INAGAGCIAA JGAGIATTIN CIGAGCITCI ATTATGGGCC CAGCATATGI NATAATTCCI TITIACACATA GGAATCIGAG GCITAGAGAA GIITIACIGAT TIACCIAATG GCACACCATA AGINCIGGGG CIAAGATITA AACICAGGIC ICCIGACITA ATTCAGATGG ICCAGCICGAT GGIAATCATA ATAATATIGI NGIIGITGIT GIIGITGITA INTATCAACA ATAGIAGIAG CIAAGICCAT ITCATGAAAC AGCICATIGG ATAGICCCAT NIGGATAATI CIGA

SEQ ID NO:835: (Length of Sequence = 422 Nucleotides)

GCTTTCTGCC TCTATAGATT TGACTATTCT GGACCTTTCA CATAAACGGA ATCATGTAAT ATATATAATA AGCAAAAGGT AACAACAACC AAGCTGGCAA TTTGGTTGAT GAATGANTAA ACAAAATGTG CTGTATCCAT ACAGTGGAAA TATTGGTGCC TACTACATGT GGATGGACCT TGGAAACATC ATGCTGAGTG AGAGAGAGCC TTGGTATTGT TTCATCTCCC CAGGAGATTC CAAGGTGCAG CCAAGGTTGA GACCCACTGA CAAGCAATGG ATATGGTTGG GTGCAGATGA AATAAGGCAG CCAGGGGCAG GAGGGATGA TCATTGAAGA TGACTGTTTT GTGGGATGCC TAGCAGGGGT GGGGGGATGA GGTATTGATA ACCAGCAACC CCAATCTTCA ACACAGCGTG GA

SEQ ID NO:836: (Length of Sequence = 408 Nucleotides)

CTCAAAAGAG TIGGCATCIC AGAAGGGAG TGTAAGINAG ACAATTGICA TIGATGATGA AGAGGACATG GAAACAAATC
AAGGGCAAGA GAAAAATTCC TCCAATTTTA TIGAACGAAG ACCTCCTGAG ACTAAAAACA GAACCAATGA TGTGGATTTC
TCCACTTCCA GITTITCAAG AAGTAAGGTA AATGCAGGAA TGGGTAATAG TGGTATCACC ACAGAACCAG ACTCTGAAAT
TCAGATTGCT AATGTTACAA CITTAGAAAC AGGTGTAAGC TCTGTGAATG ATGGCCAATT AGAAAATACT GACGGCGAG
ATATGAACTT AATGATTACA CATGTAAACA TCACTGCAGA NTACCCACTT GGGAGGATTG TCTCTAACCG GGACTGCAGT
CCAAGTAA

SEO ID NO:837: (Length of Sequence = 347 Nucleotides)

TOSCITCIGIT GCCCAGGCTG GAGTGCAGTG GCACGATCTC AGCTCACTCC AACCTCTGCC TCCTGGGTTC TAGCGATTTG CCTGCCTCAN TCTCTCAAGT AGCTGGGATT ACAGGCATGC ACCACCACTC CTGGCTAATT TTTGTATTTT NAGTAGAGGC GGGGTTTTGC CATCTTGCCT AAGCTGGTCT CGAACTCCTG GCAYCAALTG ATCCATCCAC CTTGGTCTTC CAAAGTGCTG GGATTACAGA CGTGAGCTAC TTCACCIGGC CTTGTTGGCT CTTTTTCAAA AAAAGTTTAC TNGACTCTTG CTTTATTGCA AGTCCCAGAA TGGATTTGAT TTAGGGA

- SEO ID NO:838: (Length of Sequence = 275 Nucleotides)
- AATTGCCAAG GAAAATTTTA TITTAGCITT GCATTAACAT ATTCTAAATA ATCCTITCAC TTAATGCAAT CAGATTCCTG
 TGACAAGCCA AATACTTGTT TITTTGTGTG TGTGTGTTTC CCCTTCACTT TTCATTGTAT GCCCTTCAGA AAAATCTGAG
 AAGTGGGCTT CCATTTTTGA AAAACAGGAC TTCCTTAGTA CCATAGATAC GTAGATTGCA ATTTNCCTTT TCCTGCAGCA
 TTACTGACCT TGTGAAATGA TGCCTATGGA TACGG

SEO ID NO:839: (Length of Sequence = 387 Nucleotides)

SEQ ID NO:840: (Length of Sequence = 367 Nucleotides)

GTACTAAAGC CATGCAGGAA GGAGGAAATA ATCAGTGAGC CACGGGCTGA ACTTGTGGAA AAGAAATGGA GGGCAAGGTC
ACAAACCAGT COCTAACTGC TTCTAATTTA ATGTAATCCT CACTGTTTGT CATTATTGCT TTTNATGGCC ATGAAATCTG
TTTTTCCCCCA GINCICTAGT GTAATTTGGA ATTAATTTCC CAGCTGCTTT ATTTTTTTCC TAGAAGAGTC GGGGACATTT
TCAGGATTAG TAGAGGTGTT TCTACAACAC CTTCATGCCT TCGATAGTGT GTAAGAGTTC ACCAATTGAN TTACCTTATT
CTGTTCAGAA GTAGTAACTA TGGAGTTTAA CCACTCTGGG ACATAAT

SEO ID NO:841: (Length of Sequence = 346 Nucleotides)

TGGAAAGGAA AAGCAAAAGA TTGAAGAATA AAAACATTTT GIATTTGCCA AAACTTGINC TGIAGCAGTA AGTGTGAAAC
AAGTTTGCIA CATTTTCCTT TTTGGTTTTA CTTGGTTGGG GCTTTTTTGT TTGGTTGGTT TTAAAGGATT TAGGGGATTG
GCAAGTCAGT TTGTCAGATG TCAATGAACA GAAAACCTAA GAAAAAAGGT AGCAAAAGIN CTGCTGGCCC CAGATGGATT
TINCCTTAAG TAATTTCCTA ATCATTAGIT ACAGCTCTGT GTCAAAAGAT GTACATAGAA ATTTATGCTA GATTCTTAAC
ATCTTTCCTT ACTGTGTGCA GAAATG

SEO ID NO:842: (Length of Sequence = 326 Nucleotides)

GTICTITGAA ACAAACGAGA ACAAAGACAC AACATACCAG ANTCICTGGG ACACATICAA AGCAGIGIGI AGAGGGAAAT
TIATAGCACT AAATGCCCAC AAGAGAAAGC AGGAAAGATC TAAAATTGAC ACCTAACAT CGCAATTAAA AGANCIAGAG
ANGCAAGAGC AAAGACATTC AAAAGCTAGC AGAAGGCAAG AAATAACTAA GATCAGAGCA GAACTGAAGG AGATAGAGAC
ACAAAAAAACC CITCAAAAAAA TCANIGATTC CAGGAGCTGG TITTIGAAAA GTTCAACAAA ACTGATAGNC CACTAGCAAG
ACTAAT

SEO ID NO:843: (Length of Sequence = 380 Nucleotides)

GECCTICAAA TIACAAAAAG CAATITACAT TATAGTAATA GITVATETIT ATAGTACAGG AACAAGAATG AGITAAACTA
AATATICCAA ATCAGTACAA GINATINCCT TITTITTITT TIGAGACAGG GICTCACTCT GICACCCAGG CIGICTIGCT
TIGICATCCA GECTGCAGTG CAGTGGAGTG GICACAACTC ACTGCAACTT CALLCTCCTG GGCTCAAGCA AGCCTCCCAC

CTCAGIAGCC TCCCACTCCT GATTAGCTGG GACTACAGTG AATGTGTCGC CATGCCCAGC CTAGTGGTAT TTTTAACAGA TAANTAAGAA TGGACGTAGT GGCAGAGGTG GAGTGAGANG AGAGACANGT AAAATATAGG

SEQ ID NO:844: (Length of Sequence = 257 Nucleotides)

TTTCCCTCTC GITGCCCAGG CTGGAGTGCA ATGGCGINAT CTTAGCTCAC CACAACCTCT GCCTCCCAGG TTCAAGCAAT
TCTCCTGCCT CANCCTCCCG AGTAGCTGGG ATTACAGGCA TGINCCACCA CGCCTGGCTA ATTTINTATT TAAGTAGAGA
TGGGGTTTCT CCATGTTGGT CAGTCTGGTC TCAAACTCCT GACCTCAGGT GATCTGGCCA CCTCGGCCTC CCAAAGTGCT
GGGATTACAG GTGTGAG

SEO ID NO:845: (Length of Sequence = 420 Nucleotides)

CTACACACAT CTTGCATTAC CTGGCAGTAA GCTTGGAGAG TAAGTTTTGC AGATGCAGAT CAGAAGAGAT TAGGAAGAGC
TTTGCAGATC ACCGCAAGTA TTTGTATTTC ACTCTAAATT AAACAGAAAA CCCAGGAAGG GTTTTAGGCA GATAAATGGC
ATTATTTAGT TTCTGTATTT AAGTCATCAT TTAGGTTACT GGGGGAGGCT GCCCTGAAGT GGATCAGAAG TAAAAAGGCAG
AGATACCAGC TAGGAAGCTG TTGCAGTGAG CCAGGTGAGA AGAGAGGGCC ACCTGGACCA GGTAGAAGCA GTACAGGTGA
AAAAANTCAG ACACTTCCAA ATCTTCCTCA AGATTTNATA CATTATTTGG CTGGGCACGG TGGGCTCACA CCCGTAAATC
CCAGCACTTT TGGGGAGGCC

<u>SEO ID NO:846:</u> (Length of Sequence = 215 Nucleotides)

CNCTGGGTGA CAGAGTGACC CTGTCTCAAA AAAACAGTGA TTGTTTGTAA GGAAATTATT AAAACCTTGG TTCAATATCC
AATATCTTAA CTTTAAATTT TCAAATACTT CAAAACTAGT AAGTATTACT ATGTCTAAAG CACAGTGCAG TCCAACGGAN
TATGTGAGCC ACATATATAA TTTTAACTAG GCCAGTAGTC ACATTAATAA GAAAA

SEO ID NO:847: (Length of Sequence = 266 Nucleotides)

ACACGAAGAA TCTCCTTCAT CGCCAAACAG CITTCAGAGA TAGATGCTTT GITTCCCATC GAGCATGCTA TTCCAGTGTA CTCAACATAC TGTX...'ACCTC GTGTTAGGCA CCTTTATGAA GAGATNAAGN CACTGGCATT TCAGTGGGAT TTTAAGCATT TTTAATAGCT TCATGTACAG CATGCTGCTT GGTGNACAAT CATTAATTCT NCGATATTTC TGTAGCTTGA NTGTAACCGN TTTAAGAAAG GTTCTCAAAT GGTTTG

SEO ID NO:848: (Length of Sequence = 275 Nucleotides)

CNCCIOGGIC CCCITITAAA AATIACIITI CAGCOGGCA TGGIGGCICA NGCCITGIAA TICCAGCACT TIGGGAGGCT
GAGGITGGAG GNTCACCIGA GGNCGGGAGA TIGAGATCAG CCIGACCAAC ATGAAGAAC CCCGTCTCTA CTAAAAATAC
AAAAATTAGC CGGGCGINGT GGCACATGNC TGIAATCCAG CTACTCGGGT GGCTGAAACA GAAACCACCA ACGNCTGACC
TCAGGGAGAT GTCTAAGAGC TTCTGGCATG CCTCA

SEO ID NO:849: (Length of Sequence = 318 Nucleotides)

GGAATTTINC TAGTGAGGAG TOGAGGAAGG GGGCCTGGTG GAGGAGTAGC AGCCTTINCA AAGGCCCTGA GGCAGGAATA CCTGGGAAGT GGGGCGTGC TTGTNTAAGA TGAGGCTAAA GAGGAAGGCG AGGCTTTACT TAGGAGGAAT GGGAAGCCAC TGAGTGTTAA AATTAAAAGC AGINGGGGCT GGGCACAGTG GCTTACACCT ATAATCCCAG TACTTTGGGA GGCCAAGGTG GNTGGNTCAC CTGAGGTCAA NGAGTTTNAG ACCAGCCTNG CCCAACATTG GGCTCTACTA AAAGTACAAA AATTAGCT

SEO ID NO:850: (Length of Sequence = 320 Nucleotides)

ATGICTGCCA ACTCAGGAGC AGGGCAGGAA TCAAACTITT TGGAGTTGCT ATCAAGTNCT TGATTTTNCA ATCCCAACCG TCCGCAGAAC ACTAGATGTG TGNATGTNTG CITGTGTGTG CATTTGTAGT AAAGAGGGGG TTGAGAAGTG GAAGGCAGAG NCAGGAGING GCATCTACCA NGGCATACAT NAAAGACCCT TACACCAACA CTGCCCTTCC CAGNAATGIG AGIGIAATCT GGTTTCCTAA AACCCTGGGC TGCAGTCCAG ATAGTCATGG TTAGANCAGA TGGTTGAGGA AAGGTTCAAG GCAGTAGGAT

SEO ID NO:851: (Length of Sequence = 170 Nucleotides)

CATCCAAGAT ACCAAGATAT ATGAGGGAAC ATTINNITTA ATAAAAAACA CAAAACCACA AATCCAAGAG GCTCAGNTAA CCCCAAGTAA AATATATACT AAAATACAAG NAAAAGGGAA AAAATGCATG NACACACAC TATAGGCATA TCATATTCAA ACAGITGITA

SEO ID NO:852: (Length of Sequence = 256 Nucleotides)

CAAAGIACAC ANGIGIATIT ATTACATITT GCAAGCACTC TGITCIACAT TICAAAAACG CCACCNICAA GCIGITGGCA CATTTATGIA CAAAACAGAT TAATTGIAAT GCCTGCTACA AAGCACTCTG TGAAAATACA AACTCTAATA CCAGAAATAA AAGCCAAAAG TGTCAACATC ATTACATAAG TNGAAAAGTC AGTTTINGAA ATTATCACAA ACTGTTATGN CACGGAACTG AAATACTATA ATATAG

SEO ID NO:853: (Length of Sequence = 281 Nucleotides)

GIATGINGIT TCTCTTCTCT TGCTGCTTCT AGGATATTIN ATCCTTGACT TIAGGGAGIT TGATTATNAA ATGCCTTGAG GIGATATTT INGGGITAAA TOGGCITGGN GITCTCTAAC ATTCTTATAC TTAGATATTG ATATCTCCTT CTAGGTTTGG GAAGATCICC GITGCIATIC TITTGAATAA GCITTCIACC CCATCICIT CITTATCICC TCTITACAGC AAATAAAGIT TTAGANITICO CATTITINAGG CTATTITICTA GACCCIGIAG G

SEQ ID NO:854: (Length of Sequence = 255 Nucleotides)

TCTGTCCAGG ATTATTACCA GCTAAACCAN GTAATGGAGG TCTATGCCTG ATGAAGAACA CCTGTAAAAG CTGGAAAATG TGGCTGTCTT CTCAAATGGG CAGATACCAG CACAANGATA CAAGGATTGT AAAGACTCAG AATCATGTTA CTTCCAGAAG AAACTANATA AGNTCCAACA ATGAACACAA NATAATANAA CINAAGGANA TTTGGANAAC ANTGCATAAA CAAAACAAGT TTAATGAATG ATTAG

SEO ID NO:855: (Length of Sequence = 333 Nucleotides)

ATAGCTGTGG TGGTAACCCA CCAGAGTGAG CATGCTINCT TCINAGGATA GACGTTGGGT AGTGGGATTG GGGAGAGGCA GGACAGAGGC TICCGITGIG TCTCTCIAAT TCATTGTITC TTAAAAAAGGA TITGGGCTTA CAAGITTCAA ATACTAAGAT TINATAAAGT CACATGGATT TTAAAAAATC ACICTATTGT ATGITTGAAA CATTCCATAA TTTAAATAAA AGGATTGGTA TIATATATGT NCTTGAGTTG CTATAATGIT TTACGGTTTT CCTTTGCTTC ACTITTGAAT TMINCGAGGA TCTCCTGGGG GAAGNITCAG TCG

- SEO ID NO:856: (Length of Sequence = 230 Nucleotides)
- TTINAGACAA AGTCTTGCTC TGTCACCCAG GCTGGAGTGC AGTGGCGCAA TCTCGACTCA CTGCAACCTC CACCINCTGG GITCAAGCNA TICIOCIGCC TCANCCACCC AAGIAGCIGG GACTACAGGC AOGIGGCACC AIGCCTGACT AAITTITITGT ATTITITIA GIAAAGACGG GGITICACCG TGITAGCCAG GATGGTCTCG ATCTCCTGAC CTCATGATCT

SEO ID NO:857: (Length of Sequence = 334 Nucleotides)

AAAAACAATT AGIAAAAATT AIGCATTAAG GAATTATTTA CTAGACTTTC TGGAAGTAAA AAATAAGTCA GCTGGTTTTC
CCTTTGANIT CCTATATATT AAGGCAGAAT TCTCTATACT GICCACCAAA ATCATAGTTA CAACTGTTTA CTTGAAATGA
TTTTATATACT GCATTGACCT GGCATGTTAA TATTTNCCTA TAAATATCAC CACTTATCCC CATGCCCTAA AGCAGTTTTT
TTAAACCCAT TCTTTCTTGG AGAATAATTA TAATACCTTA AATACAGAAC TTTGGGTTTC TGATCTTGCC ATAGCCATGT
AGCACAGCCA CTGA

SEO ID NO:858: (Length of Sequence = 301 Mucleotides)

GGAGAAACGC CTAATGTAGA TGATGGGTTG ATGGGTGCAG CAAACCACCA TGGCACGTGT ATACCTATGT AACAAACCTG
CACGCTCTGC ACATGTATCC CAGAACTTAA AGCATAATAA TAAAAAANTA AGAAAATGGA AATTGATTTT AAAAATTTTT
ACAATGTGCA TCAAAAGACA ACATTAAGAA AATTAACAGA NTGGAAGAAA ACATTGCAA ATAATTTATC TGATGAGGGT
TTAATATCCA GAAAATATAA AGANCTCCTA CANCTCAACA GCAANAAAAG ACAACCCNAC T

SEO ID NO:859: (Length of Sequence = 332 Nucleotides)

TGTCCTCACC CATAGAGCTA TCAGAGGGIG CCTGCNATTG GCAGACCCTT TACATTCCC TITAATAAAT CACTTCCCTG
CCAAGATCTC TGTCAAGGTT TGAGAAGTCA GAGCATTAAG TTATTINCAA TAAATGGTAT GTACATGANC ATCAGCAAGC
TCCAAGAAAT GACTCGAGGG CCTTTNACTA CTCAGAGAAT AAAGCAAAAA TGCCAGGTTT TCAGTGCTTG TCCTTTGTGC
CAGGGATTTG GACGTGTTTT TTGTTAAGIN CCAGCGTTGA GCTATGTTCC AGAAGATGGA GCCTTCCAGA AATTAATTGT
AGTGCTTGAA GG

SEO ID NO:860: (Length of Sequence = 233 Nucleotides)

AAACENTATG TGATTTTAGC ATTACAACAG TAATTCAGAA ATATCTCANN TGTTACATTG ATGTCATCAN TATTACAAAA
AAGGAAAAAA AAGTGACAGG CAACAGTGAA GAGCACCAGA GACCCAGCGC ACACCTAAAG TAGACCATGC TTCTTTCCTT
CCACTGCCAG GTTATCGTCC CGGGAAGCCC CCCACCCCCT CGCT TCCTC CTCCGCTTTC CCTAAAAAAA NNG

SEO ID NO:861: (Length of Sequence = 327 Nucleotides)

GGGCAGGIGI CAGCGCCCGI TICACCGCCA CGICGCGGAC ATGGIGATIT CAGAAAGIAT GGATATACIC TICAGAATAA
GAGGAGGCCI TGATITGGCT TITCAGCTAG CIACTCCTAA TGAAATITIN CICAAGAAGG CACTGAAACA TGINITGAGT
GACCIGICAA CTAAGCIGIC TICAAACGCC CITGIGITCA GAATITINCCA CAGITCAGIG TATATATGGC CTAGCAGIGA
CATAAACACC ATTCCTGGAG AACTGACTGA TGCTTCTGCT TGTAAGAACA TACTGCGCTT TATTCAATIT GAGCCAGAAG
AAGATAT

SEO ID NO:862: (Length of Sequence = 378 Nucleotides)

AATCAGGICC ACATTGITGI CCTGGATGCI GAGITTGCIG AGGGITTCCA AGACCAGICT CTGCGGGGAA AGGACGGCAT TGGGGCCCAG GGTGGAAAAG GGGTCCTGGG CTTCANCTGA AGGGCAAACT GCCCAGTGTA GGAGTCCGTC CAGGACAGGC AGGCAAATNC TCTCGGGGTA TGGAGATAGG TCCAACTGCC CCGAGATGTT GGCGAGTGTA ACCAAGGTGT TTTCCCGGAG CATCTCCAAG CAGTCCCACC ACCACTCCAC TTTTTTGCAG CTCACCCCTT GGGTCCTGTT CCTNCTCCTT TTCATAAGTT AGTGGTGCCT GCTTTCCGGT TCTCGGTGCT TTGTGGGTGC AGCAAGGATC AAGCTTTG

SEQ ID NO:863: (Length of Sequence = 374 Nucleotides)

TCAAATUMAT GETTTATTT CCATCTGTAA CACTAGCAGA GGAGTCCAAA GCAGACTGAT ATCCATGGAT ATAGTTTAAA
TGTAACAAAG AAAGAGTTGA ACTATGTACA TTGAAAAAAAG GAAAGACATT TTTNCATACC AACCTTTCCC TAGTTCGCAG
171/CTGAAUA GTAGAAACAA AACACATTTT TAAATCTTTC TATCAATTUA AUTUAGGAUG AAGTAACAUA ACTTTUALAA

TTAACCACTG AAGINGTCTT TAAGGACAAA ACTTAAATTT TAAAATGGGT GTTACCATAT TTNATGAGTG GACTGACTCC AAGGTTGCCT TGCTCCAAGN NTGGGCATCG TGACATTGCC GTGATGCCCA GAGG

SEO ID NO:864: (Length of Sequence = 223 Nucleotides)

AAGGGGATAG AGCAGACACT CCGCAGGINT CTTGAGATTA TCATCCGCTG AGGGTAGAGC TGAGGGTGGA AGGGGAGINA GCAGACACTC GGAAGGTGTC TINAGGCTCA GGGAGTTATC AATTATAGAA TGTTGTTGAG TTGGAGGAGG TGGCTGGTGG CCCATCCTGT TTTTTAAAGT TTCANCTGTG AGGTAGGGCC AGTAGGGCAA TCCTGAAGAA TGG

<u>SEO ID NO:865:</u> (Length of Sequence = 228 Nucleotides)

GAACCCGGGA GGCAGAGGTT GCAGTGAGCA GAGATCACAC CACTGCACTC CAGCCTGGGC AACANAGCAA GACTCCGTCT CANAATTTIN CCAAAATCTG ACGGAAAGAA AAGAAACAAA TGGTTCAGAT GGGACGGAGG GTGGGGGAGGGT GAGTAGGAAC CAGGAGGGCT GCCTGGGGTG GGGGAATAAN TIAAAAAAAG GAACGAGTTA ACAACAGC

<u>SEO ID NO:866:</u> (Length of Sequence = 328 Nucleotides)

GCACCACGIC AGAGAGGCCC CAGGCCACTG AGCCCGGGAG GAGACCCAGC CGGCCAGCCA GATGIGIGCC TGANIGCCAC
AGACTICAAG CAGITIACAA ACGAAACICA CIGITAAAAG CIGITAAAATC TCATTAAAAC AGIAGACGAG TGCITIAGAT
TCTCTGAATA TCAAATAATA TATACAGATA GACACTGAGA CATGACAGIC TAATCTAAAG CATCITIACA GATGCATTIN
CTTGAAAAAGT TAGICTICIT TTTAACICTG AATCAGTGAT AAAATTGITA ATTTGCAAAA GAGTACAGIT TTAAGCAAGA
MTAGAGTG

SEO ID NO:867: (Length of Sequence = 361 Mucleotides)

GTITCATGGC ATGINAAAAT TATGTGAAAT TCAAATTTTA GTGTCCCCAG TTCTACTGGA ACGCAGCCCC TATGTGGTTC
ATGINTTGCC TCCAGCTCCT TTCACACTGC AGCAAAGCAG GGAGTGTAAC GTACACCCCA CGGCCACGGG GCCTAAAATA
TTTCCTATCA GACCCTTAGA GAAAAATATG CCGACCTCGG ATGTGACTGA GGGTGGGGAC TTGGGTGAAT GCCGGCCAGG
AGTGACATCA AGGGTTTGAA GCAGACCCTC TGTCCAGGAG GGAGCGGAGG CAGAGCAGGG ACAGTAGTNA GGAGGCCATC
TGTGGTGACT TAGGCAAGGT GAGGAGGATG TAGGAGGCAA G

SEQ ID NO:868: (Length of Sequence = 364 Nucleotides)

AAAGCAGCCI TCAGGCIACT CTCTCTIGGN TCCTTGCTCT GGGGAAGAAC ACTCAAGCAG CTTTAGAAAA AGTCCACGTG
GCAAGGAATT GIGGICTTTT GCCAACAGCC ATGIGAGINA TCCATCTTAA GAGTGGNTCC TCCAGCCCCA GTAAAGTGTT
CAAATGACAG CAGCCCTGGC TAACATATTG ACTGCAACTT CATCAGGGAA CTTGAGCCAG AAAAACTCAG CTAACCTGCT
CCTAAACTTC TGACCCACAG AAATGGTGAG ATAATGAATG CTTGTTTTAA GCTGCTAAGN TCTGGAATAA TTTGTTATTC
AGCAGTAGNA TAACTAATAC AANGCCACCC AAGNATCATT TCCC

SEO ID NO:869: (Length of Sequence = 383 Nucleotides)

AGGGACAGAC AAGTGAGCAT CACTACCAGA GCTCTGCCTC CTGTCAGATC AGTAGGACT TTAGATTGTC ATAGGACCAT
GAACCCTGTG CATGCGAGGG ATGTGGGTTG CACACTCCTT ATGAGAATCT AATGCCTGAT GATCTGAGGT GGAACAGTTT
CATCCTGAAG CCATCCCTGT GCCCCTACCT GTGGAAAAAT TGTATTCCAT GAAACCAGTT TTTGGGGCCCA AAAAGATTGA
GGACCGCTGC TCTATAAGAA ACTATTACTG AAATAAGGTA TAAAGTCTTT ATCTTACTTA TATTTATATC CTCTATGGTG
TCCACACACA AGGTGCTTTT TACACTTAAG TTGTTAAACT AAAATATTNC TTTAAACTTT AAT

SEC ID NO:870: (Length of Sequence = 409 Nucleutides)

CAGCITICA AATCAATAG AATCATITI GCCICCNCIN ATCITACAAC TATTCICIGG AGIAGGCAGG CIGGITGAAC
TICAAGAGAA GAGGCGITCC TGAGAGCCIC CITGGIGAGC TIGCACACCI GGGGGCCAGA TGINCITIGC CCICCITGCA
AAGCCICICI AGICIGGIGC CCAGAGAATA CAGCITCAGC AGCAGCICAC TITGCITITIN AGIITAGATG AGAAAAAACA
GCAAAATAGI CCATCAAGGA CAAATTCITG CCAATGGATT INCITITIGCA AGGANGITCA CCITTGNNCC TCAAGCATCA
TCITTAAGIT GIGAATGCCI GATGGGAGGI CCAGGITGGN CIGIGGGAGG AGCINGGGGI GCNITCCAAA ACCACCIGGG
GACCAGTGG

SEO ID NO:871: (Length of Sequence = 290 Nucleotides)

TCTTTGCATT GATAGATTAG TTATTTATGC CAGINGICIC TGICTGGCTT GITTTGGTTT TNATTGCATT TGTTTGCTAG
AGATTCGTTT TAGTTTINCA ATTTCTTTCT CTGTACACCT GCCCCICCCC CACCCCACCA CTGGGTTACT ACCTCCTTTT
TGGCACTACA TGATGCCTTA AGCCCAGGNT TGCCTAAGCT TTCATAACAG ATCCCAGCAC TGCTCATCCC CAGTGGTGGA
GGINCTAAAT GGGATAACCT GATAGTGTGG GAAGGCTGGC TGGGGGTTGT

SEO ID NO:872: (Length of Sequence = 313 Nucleotides)

AAAACAAAAC AAATTTAAAA GCACTCAAAA ATAACCICAA AAAGAGACTA GIGAGIGICC CITAAGGAAA GCCCTICCIG CAGATTCCCA CAGAACTCGG CCCAGGCACT TAACCICCAT CICAGCICIG GIACAGCICA CIGCGIACAG TGIGIACCAA ACTCTTATGC CIGGNCIGCT GATAAATTCT ATTTATCICT GAACCICAAT TTATTCAAAT CTAGITATGA TATATCATAG TGCTIGIAAT TGITGTAAAA TATAGANGIA ACATACAGCA TGIGICIACA CGNITAATAA ACTGGIGCTA ATT

<u>SEO ID NO:873:</u> (Length of Sequence = 300 Nucleotides)

TAGTAAACAA GTATTACTTC AACTGATACA ATGGCTACAT GACATCAAAG TACTATAAAT NATCAAAACT ATCGTACAGA
AAAATTACAA ATTCGTTGCA AAATACATTA TACTGCTACC ATTAAGAAAA AAGTGCTTTT NGTTTTCCTTT
TTTTTTTTTT TTTTGCCAGA AAAGTATTCT TNCAATATAG AAAATCCTAC ATGTTACCCT GCATGTGGCT AGGNTATATC
ATAACGGAGT TTGTACTGAG TCCTTCTGAT TTGCTGGATG AAGGGCTGAA AAATATATTA

SEO ID NO:874: (Length of Sequence = 364 Nucleotides)

GAGTCATTGA TGCTGAGAGA TTGINAAGAA TATACTGACA GCATCCTTGT AGCTGCATCA CAGTAAATCG GACTTCTGAA
TCAAGCAGCC CAGCCTAGCA GCTGATAAGA GTGAATGTAG GTGAGAAGCA TTACCTTATT CCTGTAACAA GAGAACTGTT
TTGTGATAAG TGAAACTAGG AATGTAGAAG AAGAAATATC CTATGGCTAT TATAAAAGAN GAAGGACTTG CCTGANTGAC
TTGGTGGTGC ACCAGAAAAT AACTTTCAGA AGAATGCTTT CTGTTAAGCT GCTGCATTGT TCCTGGAGGA AATGTTATTT
CTAATGCATG TTATTTCTTC AAAAGATAGG ATAACAAAGA ATTG

SEO ID NO:875: (Length of Sequence = 341 Nucleotides)

ATCAGTCCAA TGCAGATTAG TATCACTTG CTCATAAAAG AGAGTATAAA GGTTCTTGAA GTTTTTGAAA GGAGCGGCTN
AGCTGACTGT TAAGGAAGCT ATCTTTTGTC TACAAGAAAT TTATACTTTT CCCTTCTAAA TTTCACAAAC AGAATATTAT
TAGAGACAAC AGAATACATT TACAAAAAATG GCATCAGAAA TAATTGANTA CATTGTGTAC AATATCTNCT ATTAATGAAA
TAAATGTATA TTTNATATGA TATTTGGTCT TTATGGGAAA ANTAATATAA TTNCCAATAT TCTAAGGNTG ANCAAAGNNG
GTTTACAAAAT AGCATGCAAG G

SEO ID NO:876: (Length of Sequence = 327 Nucleotides)

GTTTCANCIT GTGGGTCAAC TTCTAATATT TGATGGTGGC TACACTGTGA CAAGAAAGGT TTTTNAGCIT GTTGGGGTCA GTGGATGGGC ACAAGGGCAC CCAGTGGTGG TGCCCGGNCC AGGGAGGAGA ATACATTGTA GAATATAAGG TTTGGAAGTC AAATTATAGT AGAATGTGTA TCTAAATAGT GACTGCTTTG CLATTCAST CAAACCTGAC AAGGCTATCT CTAAGAGGCG CCAGATITCC ATGIGIGCAG TATTATAAGI TATCATGGAA CIATATGGIG GACGCAGACC TTGAGAACAA CCTAAATTAT GGGGAGA

- SEO ID NO:877: (Length of Sequence = 404 Nucleotides)
- ATTIGCTICC TGAATGITGC AGAAAACTGG TITTGITACAC TGGGGAAGGA GAGAGIGAAG ACCCTCCAGT TGGITCCTCA
 GTCAGCTCCG TICITGGIGT CGCTTTCITG CAATTITTIT CCTCCCCTGG CCCTTCCTGT GAGGGITAAA AGGGCCATCT
 CCAAGCCAGG TGGAGCCCCA ATCCCATTGA CCAAGAGGGC AAGGITATGG GTCACCTTCT CATGGAAGCC CTCTTCCTAA
 AGGAGCCCAA AGGGGACACC TGCAGAGGGC GGGCTGTGAT CTGTGTGTGA ACTTCAACAA AATCTCAGGT TAGTATTTCT
 CCAATTTCAG TTGAACCACG ATGTGGTATA CACTACAAAA TGCAGATTCT GGTGCCCCTC TCCAAGAGTC GGCCTCAGTT
 AAAA

SEO ID NO:878: (Length of Sequence = 340 Nucleotides)

TGIACOGCIG IGCIGIIGGC ACGAACACCI ICAGGGACIG GAGCIGCITI IATCCIIGGA AGAGIATICC CAGITGAAGC
IGAAAAGIAC AGCACAGIGC AGCITIGGIT CATATICAGI CATCICAGGA GAACIICAGA AGAGCITGAG TAGGCCAAAT
MITGAAGITA AGIITICCAA TAATGIGACI ICCIIAAAAGI ITIATTAAAG GGGAGGGGCA AATATIGGCA ATTAGITGGC
AGIGGCCIGI TACGGITGGG ATTGGIGGGG IGGGITTAGG TAATIGITTA GIITATGNII NGCAGATAAA CICATGCCAG
AGAACITTAA AGICITAGGA

SEO ID NO:879: (Length of Sequence = 372 Nucleotides)

GAAAAGATAA TGAAGGAATA ATGCAAAGCT GAAGGCTGTG CCAGATGTAA GAAGTGATTA TGAAGGATAA AAGAAAAGGG
CTTTCCAAGC AGGGAAGAGG CATCAGAGAG AAAACCAATT GTTGAGCCAG TATTCTGTCA CAGGGACATT TGTCTTTMTC
CTTTAATGCC CAGTAAGGGT CTTCTCAGGT TCCATTAAAC ATGCAGAATC ACAAGACCCC CCCAAAGTTA CCATGGTGCC
AACCGACTCA AAACAATACA GACAAGAAGC TCAGCTCATC AGGAAGGCTG CAGCAGGCAT ATGGGAACCA TCTTGCTCCA
CAAAGGACAG CTNAGATGGC AAAGATCCT ACAAGGGTCC ATATCCACGG GG

SEO ID NO:880: (Length of Sequence = 405 Nucleotides)

GAGCIAGGCA CCAGGCATIC TGIGAGGCCC CAGGAGITTA AGAAATGAAT TAAATATICT CCCCTGCCCT CITTGAACTG
ACTCIAACGA GGAGACTIAA GANITATITI GIAATCICIA GIIAIATITIN CIGAATTICA GAGCITAAAT ATTATACITC
AACATGAGIC ACACCITTAT TIATATGITG GIITGICICA GCIGIGITGI GGGITGGIGG AACGAGACCA CACATACATA
CACACAGAGI ACATACATGC TGITGATGIT ACACACCATAC TCACACCCCA CAAAGTGAAG CICCATGCTC ATTITGITTA
ACAAAGACTA GAGAGGCCTT GCAGACAACA GCIACCTGGA GCAGGAACAA GIGAAGCATG TITCIGAACC ATTITCICAAG
TCACA

SEO ID NO:881: (Length of Sequence = 336 Nucleotides)

GTCTTTNCAG TCAAAAGTCC TTGAAGCTGG GACCCTTTGA AAGTCTGTCA GTTACATGTT GTTGGTAGTG GCTTGTTTTG
ACCGTTTCAA AAAAAGGAAGA AAAAACCACT TAAATCATTT TTCCTTTCTC TTTTCTACTG CAAAGGCCGA CGAGATTGAA
ATGATCATGA CGGACCTTGA AAGGGCAAAC CAGAGGGCAG AGGTGGCTCA GAGAGAGGCG GAGACCTTAA GGGAACAGCT
CTCATCGGCC AATCACTCCC TCCAGCTGGC CTCACAGATC CAGAAAGGCA CCAGACGTGG AGCAGGCCAT AGAGGTGCTG
ACCCGCTCCA GCCTAG

SEO ID NO:882: (Length of Sequence = 369 Nucleotides)

TGCCATTAGC AACACTGTC AGATGAGATA ATTAAGAAAA AAAGCCAATT GAATGATGA GTGAATGANT GATTGAAAAT
CTTTCCGAAG TIATAATAAT AATTGTGATT ATTGGGGTCA AAGCAAAACC ATTTTAGTCT AAAAGATTGT ACACTATACC
AACTTTTACC CAATTTGGAA TGAAAAATTA CATTTCCAAA CCATGTAGAA ATTCTGANCT CTTTGAAATA TTTCCTTTTG
TGGGAAAGAA CCAGAAATTC TTTGTCATAT GTACCCATTT ATCTTATTIN AGTTACCCAA CCAAAAGATA AAATAATATT
CTCAAAGAGA TAATTGACTG GAGGAGTTTA AAGTGTTTAT AAATATTAG

SEO ID NO:883: (Length of Sequence = 369 Nucleotides)

CTGCCATAAG AATATCAGCC TGGGGGCAGT CCAGACGCAG CCCTTTGTCA TCCTTTCTGT TTGCCTAGTC TCAGCAGACT
GTGATCACAA GGCATTGTCT GTGGGATTTT NCCTTTCCCT TTCTTGATCT CTCTTGTGGT TCTAGGTTGT TTGGTTGTTC
ATTGTTATGG TGGCTTTTNA TTTTAACGCC CCTTGAGCCC CATGATGGCT GGTGTCACCC TGTTCCTTTA CACTGTTGGG
CCAGGTGCTG CTTGTCCTTC TTAGGGCCAT ATCAATTGCA AATATTTCCT TTTGCTCCCT TTATGAAGAT GTTCTTATAC
CCTTGCTTTT CCATATTTTT TNTGGGCCAA GCAATGCCAT CTNCTTTTA

SEO ID NO:884: (Length of Sequence = 327 Nucleotides)

AGITCATCIT TITCCAGAGG GGTCTGGGTG CCTTTAAAGG GGTGCAGGCC GAAGAAGATG GTGGCTTGGG GAAACTGGAG CTGAACTTGG ATTCAGAACT CTGAGGCACC GGGATGGGGA TGGGAATAGG GACTGGCACA GGCAAGGGGA CGATTACAGG ATACGGCACC AAGAGGGTGG CTGGTGGGAC CAGGGGGAC AAGGGGGAGC TAAAAAGGCTG TGGGGGCACA GGGGCATAGC CAGGAGGAGG CTGACAGGGT GGGGGCCCGA GAGTGCCCTG GGAGGGAAAC AAATTCTTGA GCACAGCTTC AAATGGCAAA GTGGGCT

SEO ID NO:885: (Length of Sequence = 380 Nucleotides)

CCAAAAGCTT ATCCACCATG ATCAAGTGGG CTTCATCCCT GGGATGCAAG GCTGGTTCAA TATATGCAAA TCAATAAATG
TAATCCAGCA TATAAACAGA ACCAAAGACA AAAACCACAT GATTATCTCA CTAGATGCAG AAAAGGCCTT TGACAAAATT
CAACAACCCT TCATGCTAAA AACTCTCAAT AAATTAGGTA TTGATGGGAT GTATCTCAAA ATAATAAGAN CTATCTATGA
CAAACCCACA GCCAATATCA TACTGAATGG GCAAAAACTG GAAGCATTCC CTTTGAAAAC TGGCACAAGG ACAGGGATGC
CCTCTCTCAC CACTCCTATT CAACATAGGT GTTTGGGAAAG TTCTGGGCCA GGGGCAATTT

SEO ID NO:886: (Length of Sequence = 400 Nucleotides)

GGGATGACTT TAAACGAGAG CTGGACAGTA TTACTCCAGA AGTCCTTCCT GGGTGGAAAG GAATGAGTGT TTCANACTTA
GCTGACAAGC TCTCTACTGA TGATCTGAAC TCCCTCATTG CTCATGCACA TCGTCGTATT GATCAGCTGA ACAGAGAGCT
GGCAGAACAG AAGGCCACCG AAAAGCAGCA CATCACGTTA GCCTTGGAGA AACAAAAGCT GGAAGAAAAG CGGGCATTTG
ACTCTGCAGT AGCAAAAGCA TTAGAACATC ACAGAAGTGA AATACAGGCT GAACAGGACA GAAAGATAGA AGAAGTCAGA
GATGCCATGG GAAAATGGAA ATGAGGAACC CAGCTTCGCC GACAGNAGGC TTGCCCACAC TGATTCACTT TCGGAGATGT

SEO ID NO:887: (Length of Sequence = 363 Nucleotides)

TAAAATAAAT GCTCTGGATG GGAGAATGT GGAAGTTACT TTGGAACTGG ATAATAAGTA AAGGCTGAAA GAGTACTGAT ATACATGCTA AATAAAACCA ATATTTCCCT GAATGANCTA TTCAAAAGCAA TTCTGGTGGG TGTTAGACAG GACATAGAGA CCTGGAGAAG AAGCTCCCAT TTTCATAAAG AACACAAACA ATCATGTATA GAATGTTGGT AGAAATATGA ATCGTGAACG TCAATGTAAT GAAGTCTTAG ATCGGAATAA GANAGGTTAT TAGACAAGGG AGAAAAGGTA ATCCTTGTTA TAAAGTGGCA AAGGAACTTG GCCTGAATTG TATTCATGTN CTAGTGCTTT CCT

SEO IN NO. 888 (Length of Sequence = 318 Nucleotides)

ATCHGCATG ATTAATACTA TIGGCCIGIN CCCITTATCC TCAGCIGGIT GIACAATICT TGAATGCITT CITCTICCCC TGAGGATGCT ATAGATATIG TCCIACIGIN ATCIGAAATN AGICGITTIG GAGAAGITIC TCCATCCAGA TACCTATAGA GICGGCCITT TITTITTITT TITTITTITT ATATGCAAAC NCICGCIGIA TIATTCAGGC TGATCTGAAT CICCTGGNCT TTAGGGTTTIG GACAGCTTIG GCCICTTAAA ACIGCAGGNT TACAGGCATG AGCCACAGTG CCTGGCCCATC AAGIAGCA

SEQ ID NO:889: (Length of Sequence = 349 Nucleotides)

ACAGAAATCT ACGTAGACTT CTNCCAAATG CCACATGAGA GCAGTGGCAG AATACAGAGA GACCGGCGAC CACAGCAAGG
AACTGTAACG GCCAACAGTC CTCAGGCATG CAGGCCTGGG CCAACAGCAC AACGCAGAGT CGCTTCTTCT CAGTCCAGCA
ATTAAAATGA CCATGGCAGC CAGGGTTTCA TTAGGTTACT TTCAAAAAACC ACCTTTGCTG GAAAAAATGT TTGGTAGTTT
AATCTGCATA TACGGACAGT CATGCACCAC ATAATGATGT TTAGGTCAAC GATGGACCAC ATATTCAATG GGTAGTCCCC
TAAGGTTTAT AACCAGCATA TTTTTTTACT

SEO ID NO:890: (Length of Sequence = 341 Nucleotides)

GINGIAGGE TICGIAGGIA GGECIAGIAG GIAGGGITAG TAGGIAGGEC TAGIAGGIAG GCCIAGIAGG TAGGGITCGI AGGIAGGGIT CGIAGGIAGG GITAGIAGGI AGGGITCGIA GGIAGGGITA GIAGGIAGGE TICGIAGGIA GGCCIAGIAGG GIAGGGCIAG TAGGIAGGGC TAGIAGGIAG GGITAGIAGT TAGNGCIAGT AGGIAGGECT AGIAGGIAGG GCIAGIAGGI AGGGITCGIA GGIAGNGITC GIAGGIAGGG TIAGIAGCGC GICINICCIT CITCCACCCT GGNINCITGI AAAACNITAT TITACAAGCA ATAGGAATIT G

SEO ID NO:891: (Length of Sequence = 344 Nucleotides)

GACCIGGCIG CGCACCAGGA CCGCNIGGAG CAGATCGCCG CCATTGCCCA GGAGCTCAAC GAGCIGGATT ACTACGACIC CCACAATGIC AACACCCGGI GCCAGAAGAT CIGIGACCAG TGGGACGCCC TCGGCTCTCT GACACATAGI CGCAGGGAAG CCCCTGGAGAA AACAGAGAAAG CAGCIGGAGG CCATCGACCA GCTGCACCTG GAATACGCCA AGCGCGCGGC CCCCTTCAAC AACTGGATGG AGAGCGCCAT NGAGGACCTC CAGGACATGI TCATCGICCA TACCATCGAG GAGATTGAGG GCCTGATTCT CAGCCCCATGA CCAGTICAAG TCCA

SEQ ID NO:892: (Length of Sequence = 367 Nucleotides)

CTGGGCAACA TGGTGAACCC CATCICTGCT AAAATACAAA AATTAGCTGG GTGTGGTAGT GCCTGCCTGT AATCCCAGCT
ACTCGGGAGG CTGAGGCAGG AGAATTGCTT GAACCTAGGA GGTGAGGTGG AGGTTGCAGT GAGCCAAGAT AAAAAGAGTG
AGACTCCGTC AAAAAAAAAA AAAAAAAATA TATATATATA TATATATATA TATATTTNEN CTCCAATCCC ATCTAGGTTG
CTGCAAATGC CATTATTTCA TTCTTCTTTA TGGCTGAGTA GTTTTCCACT GTGTATGTAT ACCACAGTTT ATCTTCTTGT
TGATTGATGG GCGTTTGGGC TGGTTCCACA TTGTTGCCAG TTGCAAA

SEO ID NO:893: (Length of Sequence = 220 Nucleotides)

GCAAAATAIT TATTCCAAGI TAGITATTIT ATGCAGIAGI TTCCCCCTCG AGACITGTGA TAACCACATC TITTAAATCI GIAAATAATG TTATCAAAAT AATCTTAATC TTTGAAATCI CACAAAAATT TATATTTTAC AATCCACCCT GAATATCAAG GCTGCAAGAN TAACACAACA TTTCCTATAT CCAAATATTT TACAGCTGTA CCCAAAAAGG

SEO ID NO:894: (Length of Sequence = 313 Nucleotides)

GGGATTGGGA TIGITTGGCT CIGAGGCTGT TAAGTCTGGA CIGATGCTGG AAACTAATAT CAATGTTTAA CAGGGTTGAC TGTCATTAAT GATGTGCCTA GCTGTGGGTA CAGATGCTTT GCACATTACT ACCCTCTATT CTCACAATCT TCCATGGGG ATGTATTAGA ATCCCCTTTT ATAAAGGATA AAGGTGAGGG TCAGAGAGAC TAGGAAGCCT GINCAGGGTG ACACAATACA AAGTGTCATA AATTGGGTTT GTACTCAGCC ACTCTGCTTA TTAACATCAG CAGTATOGTT AATGGGGTGA CCG

SEO ID NO:895: (Length of Sequence = 304 Nucleotides)

GGICIAGATI CAGITATGAA TGIAGGCATI AGITAAAATI AACAAGATGC AGAGIATTAA TITCITAAGA CAACA-AGIG ATTICIGIAA GITTGAGCCC TATGIGGAAA GCATTGIGGA ATCITAACCT TITTGIACAC ACTCITGIGG GACGIATCAT ATAAATGICA GCACTAAGTA ATGICITGIT TGIGGCTGAA TATTTINCGI AGATGITTIT GAAGITGACA TGACTTACGT GCATTTAAAT ATATATGCC ATCCCTTAGT TTGTAATTAA GGATTINGGA ATATGGGTTG TGGG

SEQ ID NO:896: (Length of Sequence = 337 Nucleotides)

GCAAAGIATT TCATCATATG CATGIACTGI ACCTIATTIA GCCAGCCCCA TTITGITTGG CITGIGGAGA ATTACAATAG CTGTTTTGAC TGTTGTATCA CATGCCAGGC ACTGIACTGI GTATTATCIC ATGIAATTCI CATAGITACT GCATGGTGTA GGTATTTINA TCCCCAGTTT ACAGGTAGAG AAACTGAACC CAGAGATGIT AAATAATTTG CCCAAGTTTT TTGGCTGATT ATACTGATGA AGATACTGAT ACTAGCATTC TGTTGTCAGT TATTTGCCAG ACAGAATTCT TTATTTTTTA ATACATAATA TCCATTTACT CTTGAGG

SEO ID NO:897: (Length of Sequence = 316 Nucleotides)

NATCACCINA GGICAGGAGI TCNAAACCAG CCIGGCCAAC ATGGCAAAAC CCCGINTCIA CIAAAAATAC AAAANINAGC CAGGIGIGGI GGIATGIGCC TGIAATTCCA GCIACICAGG AGGCIGAGGC AGGAGANICA CITGAACAGG GAGGIGGAGG TCGCAGIGAG CCGAGGITGC AGTGAGCCGA GATTGCACCA CIGCACTCCA GCCTGGGCGA CINAGCGAGA CCCTGCCTCA AATAAAGAAA TAAATAANIA AAGIGGGGAA GTTAGIGGIT TCTGGIGTAT TCAGAGTTGT GTACCCATCA CCCTGG

SEO ID NO:898: (Length of Sequence = 200 Nucleotides)

GAGATCIGGG GCIGGGGIAT GGATGATGGG GGGAAGGGCG GTCGCCTCTG CCACTGTCAG GGACCAGCCG GCCAACGCCC
ACCCGNAAAG GTGTCTAAAA ANTINAGCTT TTCACCCACC TGCCCCTTTC TTTCAATCCC ACGCTGTTTC CTTTCAAAGT
TCTGGGAGGA CGAACTCACC GAGGCGAGAA GTNTAACATT

SEO ID NO:899: (Length of Sequence = 264 Nucleotides)

CTCTGTAAGT TAGCGGTCAT GTTTTCAGCC CCATGCAAAG GCGCAANACN TCAGACAGCG TGGTTCTNIN AACATNAGTG
TGTGGTGCCT CCCAGGAGCA GGGATTTNAG CNAGGCTGCT GACACATAAA CACACCCCCA CCTCCAGAAG CAGAGGAGAG
GAGCCCAGGG CCAGGGCAGG TAGCTCAGCA AGGACCCCAGC ATGCTNCAGG TGGGGCCAGT AAGAGTCACT TCTCCAGCNA
GGGTCAGAGA GGAGAGAGGC AAGA

SEO ID NO:900: (Length of Sequence = 265 Nucleotides)

GCAAATGGTA AAAAACCAAG TCAGCAGAAG AAATTAGAGG AGAGACCAGT TAATAAATGT AGTGATCAAA TAAAGCTAAA
AAATACCACT GACAAAAAGA ATAATGAAAA TCGAGAGTCT GAAAAGAAAG GACAGAGAAC AAGTACATTT CAAATAAATG
GAAAAGATAA TAAACCCNAA ATATATTTGA NAGGTGAATG CTTGAAAGAA ATTTCTGAGA GTAGAGTAGT AAGTGGTAAT
GTTGAACCAA AGGTTAATAA TATAA

SEO ID NO:901: (Length of Sequence = 381 Nucleotides)

CTTCTGTGCA TATAAAAGAG AACAGTCTGG NCACTTGAAA ACAGACACCT 'ICTGGTTTTC AATGTGTTGG TCAAAGTGGC GATACAGCAA GGTTTGCAGG GTGAACACAG TGTCGGCACAT GGAACACTTA TATATATTT TREGITGTCC TATCTTGATG CCAGGATGCT GIGIGIAGGC GIGGGAATNI GIGCITGGGG CAGACITAAA CGCCATTGGA CAAATAGGAC ACTIGIAGAA GACITCACAG TGAGAACCIT GAATNIAAGA CITCAGAGCA GCCACATCAG AGIACACAAC CATTGCAAAT GCACCACATC GAAAACCAAC TCICCTCGIG TAGINCAGAC AGITCTITGI GGCGIGGGGI CINGGAAGGI G

SEO ID NO:902: (Length of Sequence = 331 Nucleotides)

GGTTGCCAGT GATCTCCTTT CTTATCACCT ATAGACAGCT TGCCTACAGG AAAAAAGAAA GCCAAACACA GACAAGCAGT ATGAGATACA ATGAGCGCCC TTGGGCCATT AAAATATGAT TGINIGCCCA AGGTCGCCTG GNCTGCAAAC AGCTCTCCAG AACCTGCAGC CAGCACAGAC CAAAGTCAGG TTTGINICCT CTTCTGTTGA TGAACAAAGG TTGATTCCAT ATCGTGGCTA TTGTGAATAG TGGCAGTAAA CATGGCAGTA TTGTATGAAA ATATNACAGA TTAGACCCTT TAAATATGTG CACTATGGNT GATCTATCAA A

SEO ID NO:903: (Length of Sequence = 389 Nucleotides)

AGCAATACTA AACATAAATG TAAATTGGGC TAAATGCTCC CAATTAAAAG ACACAGAGTG GCAAGCTAGA TAAGGAACCA
AGAGCCATTG GTATGCTGTC TTCAAGAGAC TCATCTCACA TGCAATGACA CACATAGACT CAAAATAATG AGATGGAGGA
ACATTTACCA AGCAAATAGA NAACAACAAA AAATATTTCT AATAGATTTC TGCTTTTAAT AATGAAATAT GTCAAACTCC
TATAAAAACT ATATGTAGGA AATATAAANG TTTATATATA ATTCATGTAA TGCNTAATAG TAACTGAATA GCTAGTATTG
AATAACCAAG CTTCCTTTTG TTGTTTTGNA CATTGGACAAG ATTGAACATG CTTAAAGGTA TTGGGAAGG

SEQ ID NO:904: (Length of Sequence = 285 Nucleotides)

AAATCAAGGA CCGGTTAGAT AGATGATGGG CTAGGCAGGT GGGGGAAGAC AGAGCTCACT GCCCINTGGG GTCTCTGTGG GGCCAGCCCC TNATGCCCAT GTGGCCACIN ATGCCCAGCT TCCCCCAACA CCCCANCACA GGCCCAGGTC AATATTACAA AAGTGAACAA ATGCAACCTG TTTCTGCTTT NACAAATGAC ATGTCTCCAT CCCCGGCCAG CAGGGGTAGG GGAGGNCGGT TGAAAGTGNC ACTCCGGTTA AAAAGGCAAC AACTTTTATA AAATG

SEO ID NO:905: (Length of Sequence = 374 Nucleotides)

GAAGCAAAAA GITGAACCIT TTAAAGTGCT GAACACAAAT CCAAATTCGA ATGGITCAAG CAGCCGTGAA ATCGCTCTTC
ATAAAGTGGG CITAATTCTC TAGITTAAGT TCTTTTGATG GAATGAATTA ATTAATGTGT CAGGIGGCIT ATTITGTGGAT
GCCATGATTG ATGATGTTCA TTTTAAGCTC TTACCTATAG TACAAGTACA TGATGCTACT GAATATTTTT TCCACTTGGA
AACTGTGAGC TGGGTTGTTG CATTAAAACA CACATACANA CANAATCANN AAACACTGCG GACTTTTCAC TCAAGCTGGG
TCTTTTCTTC CCCAGTGGTA AGGGCAAATC CTGGCCTANC TAACCAACAC CCAC

SEO ID NO:906: (Length of Sequence = 375 Nucleotides)

CTGACTGAAA GGCTCTTTCC AGCTCCAACA CATGAAGGIT CCATAATTIT CCCCAAATGI CTGCCGCTCI GAAAACITCA ACTATCTIAA TATTIGIGAC ATTIATGCCI GIGIATGGCA ATCIGATGGI AAAAGGAGCC ATATGIAAAT AATAACIGAA ACTITGICAA AATAATGITA AGGAAACATA ATTAGCAAAG CAATATATAA TINCAAGTCC ACTGATTTAG AGAATCAGAA GIAACANTTA GAATCAGAA TAACAACTAT CTGGCAGGGA TGGAAAAATG AGAGCAGATA TAAAAGGIGT ACCCCAACCC CTGACCCCAC TGCCCATTIG GGIGIGCACT ATGINITICC AATATTAATA TCTTT

SEO ID NO:907: (Length of Sequence = 390 Nucleotides)

GIGCIGACIT CAGCAGCCCI CIGAAAGGCC CCITCCATAA GCIGGGAAAG TATGATCATG GITTCATCAT CCITGITGGI TATTACITCA AGGITGACCA ATCIGAAAGC TCIGIGIGAA GAAGGGGACT GAGIGGCIGI GAATGATGAG ACCGITGITI AAAAGCCAGG CITTACCATAG GGICCGGAAG AAGCAACCIC ARTUCIGIGC TITACCATAG CACCACCIGC AGGIATCCAG GAATAGAGAA CCCAGCTGAG CGACTCATGC TINACCAAAA ATACCCAGAG CAGTGTGTCT CTACCTTTTT AAGCCCATGC TCACTAGTGG GGAAAACAAT TTTACCCCCC TGTATTTAAA TATGGGGATT TCAAGGCAAA CAAAAGCATT

SEO ID NO:908: (Length of Sequence = 207 Mucleotides)

CTIGCATACA GGIGGIAAGI TATTACATTA TITCINCCIC CIGICIACCI GCAGITGGIT TIATGAGGGG CGITAGIACA

CTICCCAAAG GGCTIGCCCG CAGGITNAGA GGIGCACATT GAACICCCIC ACCAGGCAGA TGGGAAGIGI GGCCATGAGA

GAGAGCTICA GGGGNCCING GNITATNACA TCGCTGGGCC AGGANAT

SEQ ID NO:909: (Length of Sequence = 339 Nucleotides)

GCAAGAGAAC CIGATATAAT ATCTATAAAT TITGATTCCC TGGGGTATAA CAAGTAAATA ATTTTTAAAT GGIGCTTAGC

AAGATTGGIT CATGGNAAAT GAAGCAATTA TGGCITGANT TTATATGTAC AATATTTATT GICTTAATTI TAATTTAAAA

CGAATGACAT GICTCTTTT TTAAAAAAAG TCTTCTTTTA AAGATCTTGT AGTTGATGTG ATGAGCTATG CACTGCTAAA

TATTTATCCA CACATAAATA TTTGANAAGG AATATGGNAT AGTCATGGGA TGTAGTTTCA TCTCAGTGCT CCATGGAGGG

AGTGTTTTCA CCCCCCCCT

SEO ID NO:910: (Length of Sequence = 372 Nucleotides)

CTCAACTGCC ACTCACCTAT CTACCATCCA CTACCCANTN ACCACCCACC ATGACCCACC ATTTGCCATC TACCCATCCA

TCCATTCTAT AAATAATTAG TAAGCACTTA ATGCATGCTA GGTATTATTT TAGGCACCAG TAAGACAATC ATGGCAAAAA

AAGACAGACA ACCCCCGACC CTCCCATCCT CAGGGAGCTC TATTCCAGTG AGAACAATCA ATGTGCTAGA TTGTGAAGGT

CATCAGTGCT TGCTGCCCGT GTAAGACTGA GGTTCCCAGG CCCGAGGACC AGNCTGGGCC AGGGCTTCCC AGGGGTCINC

T. LOGGGGA CTCTCAGGAG TCCAGGTGCT GCCCCTTAGC TNAGCACCTG GG

SEO ID NO:911: (Length of Sequence = 377 Nucleotides)

GAACTICAAA AAAAAAAAA AAGAGGAGIC ATAATAAATA TITNACIGIC TAGICAACCI AATITATGAA GCCIGATTAT

CTAGCINAGC CICCGGAGAT TGCTACCGGA AATCTCCCCA GATGTTCCCC CTTCTAACCI AACINTCCAC TGINIGGCAG

GAAGGCAGCC GGGCATCTGC ATTCCGGAAG CCCAGCTGCT TGGGAAGAGA GAGGGAGCGG CCTGCACGIN ACTCAACAGC

CCTGCCTGCT AACCAGTTAA CCAGTTCTCA GTTGGGTTCA CGGACCCATG AGCGACCCAG CTTTCTTCCC CTCAGGTTGA

TATTGTGCTC CAAGCTNGGG GATGCCCCGG GGGACTATGT GGAGGGAGAG TTCCTTA

SEQ ID NO:912: (Length of Sequence = 370 Nucleotides)

ACAATCTACT TGCTACAGAA TCAGGATGTA TTINCCTATT TATAATAAAC TACAGAAGGT AGATTTCAAA GGTAATGGCT
GTTATGGAAA CCTACTTGAG GTTGTCTGCT AAAACCAACT CAGTGTGCAA AGCGAAATAC ATTINCTACT TCAATAGCTC
CTCATACTGC ATCTGTCTGT AGAGTTTATT TCAGTAAAAC TGTTTACTAT TTCATGATGA GTAGCTAGAA TTAAAGCATT
AAGTAGCTTG AGAAAATAAT CTATATAAAT CTTTATATCC TACATATGGC TATAAAAAATA AATTTATAAT TTTTAAAAAATT
GTTTTAAATA AACATTTATT TTTTACCCTA CCAAAGTAAA GGGTATACAG

SEO ID NO:913: (Length of Sequence = 313 Nucleotides)

GTATCTGGIT GCCACATCCA AGAAGAACGC GTGCNINICG CTGGTCTTIN CTTTCCTCTA TAAGGTGGTG CAGGINITIT

CCGAGTACTI CAAGGAGCTG GAGGAGGAGA GCATCCGGGA CAACTTINIT ATCATCTACG AGCTGCTGGA CGAGCTCATG

GACTTCGGCT ACCCCCAGAC CACCGACCA CAGGACTCCTC AGGAGTACAT CACTCAGGAA GGCCACAAGC TGGAAACAGG

GGCCCCGGGG CCACCAGCCA CCGINACCAA CGCGGTGTCC TGGNGGINCG AAGGCATCAA GTATCGGAAG AAT

SEO ID NO:914: (Length of Sequence = 389 Nucleotides)

TTACAGGCGC CTGCCACCAT GCCCGGCTAA TITTNAGTAG AGATGAGGIT TCACCATGIT GGCCAGGCTG GTCTCAAACT CCTGACCTCT GGTGATCTGC CCACCTCAGC CTCCCAAAGT GTTGGGATTA CAGGCGTGAG CGACCGTGCC TGGCCTTCTC CACTGTTTTC ATAGTGAAGA AAGGACACCC AAATTTTGAT CTGGTTCAGC TATTCACTAT TCTATCCTGT GTGGTCTTAA GCAAGTTACA TAACTTGCCT ATATCTCAGT TTACTTAGCT ATAATATAAA TTAAATTGGT CAAATGTTCT CTAAAGTCTT ACTAGTTACC AGTGTTCCAT GGGCCCAACA GCATCTACAT TACCTGAGGA GGCTGGTAGG AAATGCAGG

* SEO ID NO:915: (Length of Sequence = 328 Nucleotides)

CNCCAGCAGA TITINATIAG ATGGAAGATA ACAAGCATTA CCNCATAGGI AAGIGGIAAG AAATGGCAAG TACAGCCAAG CCACAGAGGA GIGAGGACAT TACTGGCTAT GGGAATGGT ACTTATGAAA TCTAAGGGIT GGGTCTCCIG ATGAACTCTA ACTACCCAGT AAGCTCTTCT CTTTGGCACT CAATATGACC NCTGCTGGCA TGAAAGGGNC TACAGTAGCT ACTTTCAACT TGGCCAACAG TTCTTCCAGT TCTGGTCGAG CTTTGAATCG TCCCTTTGAA GTCTTTCTTC ACNTGGTGCT CCTTCAACTT GACAAGTC

SEO ID NO:916: (Length of Sequence = 365 Nucleotides)

CAACTICAAG GIGCIGCAAG AGCITICAAG AAGAIGGGIG TIGACAAAAT CATICCIGIA GAGAAATIAG TGAAAGGAAA ATTCCAAGAT AATTTINAGI TIATICAGIG GITTAAGAAA TINITIGACG CAAACTAIGA TGGAAAGGAT TACAACCCIC TNCIGGCGGG GCAGGGCCAG GACGITAGGGC CACCICCITAA CCCAGITCCA CAGAGGACGI CCCCCACAGG CCCAAAAAAAC ATGCAGACCT CIGGCCGGCT GAGCAATGIG GCCCCCCCCT GCATTCTCCG GAAGANICCT CCATCAGCCC GAAATGGCGG CCCATGAGACCA CAGCT

SEO ID NO:917: (Length of Sequence = 400 Mucleotides)

SEQ ID NO:918: (Length of Sequence = 348 Nucleotides)

CTATTGCACA TGGIAACTCT GICATACATC TATAAAGCCT AGTAGCTGTA TTGGGIGAGA TGAAAAAAAC TGCITATATT
CCACAGCAAC ATAATTACAA ATAAGTTTIA ACCIATTAAA GIACAGAGTC TCTCTCATCA CTTTCAAAGC AGGACCCTAC
TTACCAATAA TTCATAGCAT ACCICCCCTT ATTTTAAAAC TCATATGATA GCTGATTTCC TAACTGTAGC AATCAGGATT
CTTAGAAAGA TTCGAAACTG AATTTAGCTA ACTAAGGAAG CGGATTTCAT TAAAAATATT GGGTTAGTTT ACAGGAATCA
GTAGTGGAGG AACCAGGGTT GCATAAAA

SEO ID NO:919: (Length of Sequence = 345 Nucleotides)

GCGATGACIT TAAACGAGAG CTGGACAGTA TTACTCCAGA AGTCCTTCCT GGGTGGAAAG GAATGAGTGT TTCANACTTA
GCTGACAAGC TCTCTACTGA TGATCTGAAC TCCCTCATTG CTCATGCACA TCGTCGTATT GATCAGCTGA ACAGAGAGCT
GGCAGAACAG AAGGCCACCG AAAAGCAGCA CATCACGTTA GCCTTGGAGA AACAAAAGCT GGAAGAAAAG CGGGCATTTG
ACTCTGCAGT AGCAAAAGCA TTAGAACATC ACAGAAGTGA AATNCAGGCT TGAACAGGAC AGAAAAGATA GAAGGAAGTC
AGAGGATNCC ATGGGAAAAT GAAAT

SEO ID NO:920: (Length of Sequence = 299 Nucleotides)

CCCAGGTACT CAGGGAAGGG GCAGGAGAAC CACTTGAGCC AAGGAGTTCA AGGCTGCAGT GAGCTGTGAT CACACCACTG
CATTCCAGCC AGGACAACAG AGTGACATCC TGTCTCAAAA ATAAATAANT TTTTTAATGA TGAAACTAAC TAAGGTACTG
AGGAGGTAAG ATATTCCCC ACGGTAAGTC ATTCAGAAAC TAAATGTGAA AAACCAAAAG AAGCCTCTGG GGTTAGTATT
CCCAGTCTCC TTGTCTGCCC AGGACCCCAC ATTTGTGTAA GTTGCTAATT GCACAAGGG

SEQ ID NO:921: (Length of Sequence = 234 Nucleotides)

ATGAAGCAGA GGCAACCAAC AGAAATTGAC ATCAGAAACT CTGCTGGNTC CCCACCAGCA TGCTACCGAT GANTCCTGCT CTCTTTCAGA TGAAATTTTA TTTTTTTTNCC AATAAGGCCA GCCCTACCCT GGAATCTGGA ACCANTTCTG GCCCAGGGTA GAAAGGCTAC CAAGCACCTA TGGTAGAAGC CCTGGTGTCC AGGNATGCCT TGGNCCTTAT TATTGACCTT CTCT

SEO ID NO:922: (Length of Sequence = 328 Nucleotides)

TAGCAGGETT ACTGGCCTTG GCTGCGGCCA AGGGAAAACT CTGCAGGCCC TATTACTTGG CGGCCTTTAA CTCTTATAGA
ATTGGGAGAG AACACTGACA AAAGCGAGGA CATGATTITN CGGTTACAAA TNATTTTCCT TGCTTGCTTT CTTCTCACCC
TTTTNAATTT TCCTTTTCTN CTTTTCCTGT CTATCTTACC TTCCCTCCGT GATCCCTGCC AGCCCCTCCT TTCTTATTAT
AGCTGATCAT GGCAGTATTG TTTTTNCTG GGTAAAAATC AGAGTGGGAT TTAGAGAAAG CTTAGCAGGC CTAGCATGAG
GGCCTTAG

SEO ID NO:923: (Length of Sequence = 371 Nucleotides)

CAGGAAACCT ACTGTGAAAA TGCAGAAAAA CAACAG:CAA AATTGATTGT TGACTCAATA TGATATATAG TTCAAATGTA
AACAAATGCT TGINAGCATT CCACATCACT GAAGGAAAAA AAGTAAGTTA TTATTTCCAA TGITGGGAGT TAGGTTGCTA
TAAGCTTATG ANCACACACT TTCAGTGAAT TTATGTAGAA TCGGAAGCAC TTCATTCTCC CCTCACCACA CATCACCCCC
TTGCTCCTCC TCGACACGTG CAAAATGATA GGGCATGGTA GGGGTTGTAG TGAAATNGAG AAGGCATGCC CCATCTCAAG
AAACAGGGTG GACCAGCCAC AGCTTTCAGC TCCANTTTGT GATACAGGAA T

SEO ID NO:924: (Length of Sequence = ucleotides)

AGENECISCI TITITITICAT ACCITTACIT TINAS AGENECISCI TITCIGGASC CGACTGAGGG ACTGGAGAAG GCTACGGGG TCCTCGCCCT GCCAGGGCAA TCCTT. CTCTTNATCA TITGGTTATIG CAAATCGCGG TAAAGTITTTT CCGAAGGGGG TGCTGGCTCC TCTTGGCAGC TCTTCTTTTT GACTTTGGGC ACCAGGGCTG CTCATACCTG CAGCCTTTTC GGCCCTCTNG GCCCCCAGGCC GTCCGGCCTC CCGAAGCACT GCCATGGCCC GGAATAGCAG CCCCNGAGCA AGG

SEO ID NO:925: (Length of Sequence = 317 Nucleotides)

AATGCTTTAT GATCAACTTG CCATAGGACT GATGGATTAA CCAGTGTTCG GCTTTATTTG AAGTCTATGC CCTGCACAGC
TCTTGTATGT ATTINAGATG CTAGAAGTTT TTINAGCATG TNATGTGTGA TTCTTGTTTG AATTCTAGGAN ACCTTGTCCA
ACTTGGTTCT TTTTCAAGGT TGTTTTGGGT ATTCTGGGTC CCTTGCTTTT CCATATCNAT TINAGGATCA GCTTGTCAAT
ATCTGCAAAA AAAAAATCAG CTATATTTTG ATAGAGNTTT GTATTGCATC TTTAGGANTG GTTTGTTGAG TATTGCC

SEQ ID NO:926: (Length of Sequence = 247 Nucleotides)

GITATTCATA CCACAGCATT TAAAAAGCAA TCCGCAAGIN ATAAAAAAAA AAAAAAAAA ATGATGIGAC ATATCCATTG CCTGANITGC CTCTTTTGIA AGCCAGINIT GGGATTATAG CAGAGGAGTA GCAGAAATAA NTATATTCAG ACACAAACAT ATAGATATAA TAATATCCAA CCNCTTTATA TGATTTAGGG TCTCGTTAAA A'GGGTTACCA TTTGCTTCTC CTAAAANTTA TATAAAT

SEO ID NO:927: (Length of Sequence = 286 Mucleotides)

GGCTGTCATG AGAATCACTT GAACCCGGGA GGCGGAGGTT GCAGTGAGCT GAGATCATGG CACTGCACCC TAGCCTAGGT

GACACAGCAC AAAAAAAANC AATGTTCCAC AAGTCAAAAA TTGINTTCAG GGAGTAGAAA AGTAGTAGGC TAGGTATCAA

AGGGTATGAA TGACTAAGTT CCTTCIATAA TATATTGACT ATAGGTTAGG AGATACACTT TCAGTTCCTG TTTTINGTAG

ATCTCCCAAT GATCIGTCAT TTAAGAGTAC ACACGATGAG TGGAAA

SEO ID NO:928: (Length of Sequence = 349 Mucleotides)

CTIGITIAAC CAGIATTIAT TGCACATGGI TITGITATCT ATTGCATGIG GIAAATTACC CCATACTTIG CTICITAAAG

CATTAGACAT TICIGIAGGI TAAGAATTCA GAAGCAGCIT AGCIGAGCAG TICITGCTCA AGGICTGCA TGAGGITGCA

GICAAGGAGC TGGCCAGGGC TGCAGICATC TGAAGGCCTG ATTGGGGCTG GAAGACTCCC TTTCCAGATG GCTCCCTCAC

AGGCTTGGCA TGTCAAAGCT GGATTGTTGG CAGGGGACCT CCATTCTTCC CCACATGGGC ATCTCCATAG GCTGTTTGAC

ATGGCAGATN GCTTCCTCCA GCAACTGGG

SEO ID NO:929: (Length of Sequence = 395 Nucleotides)

AGAGGAGGCA GCAGCCACCC CCAAGAAGAC TGTACCTAAA AAGCAAGTTG TGGCCAAGGC CCCAGTGAAA GCAGCTACCA

CCCCTACCCG GAAGAGTTCT AGCAGTGAGG ATTCCTCCAG TGACGAGGAA GAGGAGCAAA AAAAACCCAT GAAAAATAAA

CCAGGTCCCT ACAGTTCAGT CCCCCCGCCT TCTGCTCCCC CACCAAAGAA GTCTCTGGGA ACCCAGCCTC CCAAGAAGGC

TGTGGAGAAG CAGCAGCCTN TGGAAAGCAG TTAAGACAGC AGTGATGAGT CTGATTCAAG TTCTGAAGAA GAGGAAGGAA

ACCCCCAACT AAGGGCAGTA GTCTCTAAAG CAACCACTAA ACCACCTTCA GCAAAGAAG CAGCAGAGGA CTCTT

SEO ID NO:930: (Length of Sequence = 214 Mucleotides)

ATCCAACAAT GACAACICCT CITCGGACAA TATTGGCACT CCATTCAAAC CITGITTCAG GICAGICCGC ACITCATCAT

CTCCCGAATTT GICCAAAACA TACTGTAGCT CAAGIACAGT TTTTAAACGT TTCTGTNCAG CITCITCTCT CATAAGCTGC

TCCCGACGTG CTGTCTTCTT NATTGTTTTC TGAATATCTT GACTTAGTGC CATG

SEO ID NO:931: (Length of Sequence = 245 Nucleotides)

GAAAGINITC ACAAACATGA TGCTIATCIA ATAAAATATC ACTGAGCAAT AAGGAGAAAT ATITIAAATA GATTTGAAGT

TGTGAACAAA TAATTTAGAG TCCAAAGAGG ANAAAGANAA TTAACTCTGT TTTINATCCC TAGAACTCAG AAACTTTACT

GGATTGGTCA ACAAAGACAA ACTTTTTATT GTATAAAACA GTAGANTTCA TGGAAGGGAT AATNCTTTTG GAACAGGCTT

CTCGG

SEO ID NO:932: (Length of Sequence = 303 Nucleotides)

CATATIGGG GCCCAATATA AAGCAAAGCT GGAAGAAGGG ATGATCCATG TATTINIGGG GATGGGATAT GGACAGGGAA

ATAGIGTICC AACTCCATGC TGAGIGTIGI TITGAATTGT AATGIGAAGT TGCCACCATA CCAGGGCTAT GACTGINTAC

GATGICTCAC CCTTGTAGGC TAGTAGCTTT GCAGIGGGAA AAGATGACAG GGCCACTTGT CCAGGGCATT CAGGTAATAA

AGTCCCTGAG CTCCAAGTTG CTAGATCTAA GGAAGTATTT TTCCCTTCAT GTCAAAGATG GGG

SEO ID NO:933: (Length of Sequence = 186 Mucleotides)

CTCTTTTGGG CTG-ITCANA TCTCCGGGGA ATTGAAAGCA GTGATCTCTC AGGTGCTAAC CGGNATAGTA TTAGAAGACT

CCAATATCTT GCACCTGTG GGACTTACTG TATTL-TCTT TGTTTTGTTT CATTTGCTTT TGGGTTCTTG GTCATGAGGT

TTTGCCTAAG CCA-IGTCTT CAAGGG

SEO ID NO:934: (Length of Sequence = 336 Nucleotides)

GOGRARACGI ATCAGCACAT GARATACCIT GIAACIATIT CATITATATA ATTIGCTACG TGITCITIGC AACATAGIGA

AAAATAATCA TGICTGATGI TIAGTAGGCA CATAATAAAT AGIAATGGAA TGAATGGITG TATATITAGA GAGCCATGCT

GAAAGGITAA ATAGCAAAAT ATGACTACIT GGAGAATAAT GITAAATTGI CAAGGAGAGI AGIGTTATAT GAATACTCAG

ATGGATGGAT ATATAGANAA TGAGAAAAGC GACAGAAGGA ACTTAAAGAG NITITAAAAA TAGCTTTGIC TAAAGATTAA

AAATTAAAGG TICTAA

SEO ID NO:935: (Length of Sequence = 383 Nucleotides)

AGGITAAGAAA ACTGCTGAGI GGGCTCCTIG TACCAGCACC AACCAGCAGC CCITGACAGC ATAGATGGGA TGAGTGTAAG

GGCTATCCTT AGCATAAGGG AAAGACGGIT ATAAGCTGAG AAGATTGAAA GAAGAATGGA GCCACAAAGA GAATAGCATA

AATAACAAGA AGGAAACATG AAGAACAAGC ACTTAAGNTA TTAACTTTCA GTCTTTCTCC ATTTCTTGAT GTCTAATGAG

GCAAAAATAAC TGGGCAAGGA CCACCAAGAT GAAGAAGTTA AATAAAATGT CACAATGAAA TTNAGGTGCA ATAATACAAC

TGTTGACTGA CTTTCCAAAA CCACGGTGAT CGGTAGAGTA TCATCAATGT TACCGAGGAT TTT

SEO ID NO:936: (Length of Sequence = 204 Nucleotides)

GAAGCTGTGC CACCCTTCIN AACTITNATG AGCTGCCTNA GCCGCCAGCC ACCTTCTGIN ACCCAGAGGA AGTGGAAGGG

GAGCCCCTGG ATGCCCCCCA NACCCCAGACT CTGCCCTCAG CCCTTGAGGA GCTGGAGCAA GAGCAGGAGC CGGAGCCCCA

CCTGCTAACC AATNGCGAGA CCACCCAGAA GGAGGGGACC CAGG

SEO ID NO:937: (Length of Sequence = 386 Nucleotides)

CTAACTAAAT AAGGGTTGCC AGATAAAGTA CAGAAGGCCC AGTTAAACTT GAAATGCATA TGANCAAGAA ATATATTTNA
GTATGANTAT GTCTCATGCA ATATTTGGGA CATAATTATG CTAAAGAAAG TATTCACAGT TTTNCCAACA TTCAAATTGG
AATGAGTGTC CTGTATTTN ATTTGCTAAA ATGGGCAACC CTAAGCTGGT ATCTCTACAG TTACATACAC TTACCAACCC
CACCCATTCA TACTGGTCCA AGTTACACCC CAAAAGAGGG CAGAAACAGA ATCTGAACAA GCTCAAGTTT NGAGGGCAAA
AATGTTTCAT TCTGCCTTCT GGATTNCTGT ATGAAGACTT TTGTTGTGAA AGATATGAAT AGAACC

SEO ID NO:938: (Length of Sequence = 349 Nucleotides)

GACACTITCA GAATTAAGAA GCCTTGCCCT CTTTGCGTGT CTTCACAATT GINITAAGTC TATTATAGTA TTCATTITAG

TTTGAAAGCA ATAAATACAA TATTAGTACA AGCACACTGT CAAGAAATCC CTAGAATATG GCTCCTCTGA AGGTTGACAT

GGGTCTGCCT CGCATGTATC TTTTCATCTC CAGCATCCAG ATCAGAGTCA ACAACAACAA CTCTACAAAT ATCAGGCTTC

TTGGTGGGAAA GAAATCTGGA CATTTTTNCT ATGAAAAAAA AGTTAGGTTA CATGGCATTA ATATTTTTGC TAGACTTAAC

CTACAGAAAA TGTTTCAAGC TTATAAAAA

SEO ID NO:939: (Length of Sequence = 374 Nucleotides)

GRAATAAAGC CICACAAGAA ATAAGGIGCT TATGGIGITA AGITACAATG GAAAATAATC AATGGCATTT GTATGCATGC

TGCATGIGIG ATGIAGATCA GTICATAGGA GATGGGGCAA CAAATAAATA TCACCATGGG GATGTGATCA TCAAAACCCA

GGCTGTGGAA AACTGTCAGT CAAGITTCIT CAACATATTG CAAGAAAAAT ATGATGGCTT GAAAATCTAT AGATGAAGCA

ATTTAACAAA CCTACCAATC TCATTTAATC TTGATTACTT TTAAAAAAAG ATTAAAAAGA TGACAGAGAA AGGGTTTAAA

AATTTGTAAG ACACGCTGG ACGGTGGGC TCACACCTGT AAATCCAGCA CTTT

SEQ ID NO:940: (Length of Sequence = 385 Nucleotides)

GTAATCCCAG CTACTTGGGA GGCTGAGGCA TGAGAATTTC TTGAACCCGG GAGGCGGAGG TTGCAGTGAG CAGAGATCAC GCCACTGCAC TCCAGCCTGG GCAACAGAGC GAGACCCTGT NTCAAAAACA ACAAAATAAA TTTCCTTTTA ACATCTGINC CAAAAATGAG ATAAGCGTTA TCAGGGCAAG TCCATCCTCA TCACTCTTTC CCTCCCCACT GCCCTCTCCA CGATGCCCAG CTGATCAAAA GTCATTTTTA CTCATAAGAC CAAAGTATCA TGGGATACTG TGCAGTINGA GAGCAGGTTG ANCATCAGAA ATAATTGCTG ACAATAAAGT AAAAGATGGG AGAAAAGCAA GGCCNATTGT ATATAATACA GCTTC

SEO ID NO:941: (Length of Sequence = 406 Mucleotides)

GGTAACAGGT TITTACCAAC AATTGCTTGT AGCTAATGTA GAACATACTT GAGAAAATGG CTTCTGTGAA AGACCAGTTA

GTACCAAAAAT AATCTGGCCC AGAAAAATAG CCACCATTCT TGACTACATT AATAGAAATA GAATAACCCC CAAAGGGAGA

TGAGAAGCAT TCTAAAAGTGC ACTGATCATG AGTTTCTATG TGATGATTTG TGTCCATTTG GAGCTCCAGT GCTTTAAAGC

TGAAAATGAAT CCTGGCCTTT CACCACCCTC CCTGCCCATA GTATGGTATA TCCTCTTATT CCTTCCCTCT TAGCTTACTG

AGAGTGTAAT TTCCAACCAG TTAAGGCCAA AGAGGACTAT TTTCTAGGAA AGGAGAGAG GATGAATTAG CAGTTAATGG

AGGAGT

SEO ID NO:942: (Length of Sequence = 296 Nucleotides)

GATGGCTCAT GCTAGTTCAG CAAATATTGG GCCCTTCCTG GAGAAGAGAG GCTGTATCTC CATGCCAGAG CAGAAGTCAG

CATCCGGTAT TGTAGCTGTC CCTTTCAGCG AATGGCTCCT TGGAAGCAAA CCTGCCAMTG GTTATCAAGC TCCTTACATA

CCCAGCACCG ACCCCCAGGA CTGGCTTACC CAAAAGCAGA CCTTGGWGAA CAGTCAGACT TCTTCCAGAG CCTGCAATTT

CTTCAATAAT GTCGGGGGAA ACCTAAAGGG CTTAGAAAAC TTGGCTCCTC AAGAGT

SEO ID NO:943: (Length of Sequence = 223 Nucleotides)
GIGCCATTAC AACTITINCIG TAACCCIGAA ATTGIGICAA AGIGAAAAIT TITTAAATGA GATTATAAGA GCATAATCAA
ATTGGAATTT CCITAGGATA CCAGAGAATC ATTINCITCI CAGGIAAAGG ANITITCCIT TINGTAGTCC AGAGCIATAC
ATGATTAAGA AANIGITCAG NCCAGGAAGA TGACATCICI GCTAACCIAA TCGATTATCA TGG

SEO ID NO:944: (Length of Sequence = 327 Nucleotides)

CCAGGCACTC AGGCTGGCTG TCCCTTINNT CCTCCTGCCC ACCCCATCCA CTCTGAGCAT CAATGCAGCC GGCCAGTTGC

AGGCAACCAG GCAGCACCCT GGCTGCCCAG GCAGGCTAAG AGGCCCCCAC CCACTCCCCC CTCCTTTGCC AGTGGAAAAG

CTTGCCGTAG GCATAGCTTT CCCAGCCTTC CCTGCTTCAN AGGCAGGAGC ATGGCACTCT GGGAGTTGTA GTGCTCATAA

CACTCAGGCG ATCCCTTGTG CAAATAACTG GAGGAGAGA CTATGGTATT GGGGAAGAGA AATTNAGGAA TAAGCAAGGA

GTTGGCT

SEO ID NO:945: (Length of Sequence = 222 Mucleotides)

CTIAAACAAT AAATACACCI GAGITAGITI TOCAAACCII TOCIOCIGAT TAAAATGCCCI TAAAACTIAA ATCICITGIT

ATCITCAGIT GIGATCIAGI COCAAGIGGA AATTACGITI AGCITTAAAA CCATGAATTI AAAGCTCAAG CCIGIAGCIG

GCIGCCIAGG CANITITATGA TTAGITTCAC AGAATAGCAC CCACTGGCIA CACAGGNCCC AG

SEO ID NO:946: (Length of Sequence = 286 Nucleotides)

GCICICICIA COCCCICATO TAGGITAGIN TATAGCICAT TIATTIAGGG GIGATGITAA AAAATIGAAT GCCCITAATG

GCAAGGGAAC CAACCAATCA ATGIGGATGC CACAACTITI TCCCCIGITG ACTGITGINA TIXGITATGGA ACTATITITT

TUTTICICCCA GCITITATIT CAGGITCAAG GGATACATAT GCACGITTGI NACATGGGIA AATTGCATAT TGIAGGGGIT

TAGIATACAG GITATTICAT CACCCAGGNA ATAAGCGIAG TACCTG

TTATAACATT TIGAAGAAAA TCTTTAAAAA TNTTTGTTTA CACAGAAAAT AATCTTAGAA A

SEO ID NO:954: (Length of Sequence = 217 Nucleotides)

AGAGCITAAA AATAGIGAAG TCTTTATAAG TAATTITTAA AAATTTAAAC TAGGACCATA AATTICTAAA CTATGAGATA AATGANCAAG AAAACAAACA GGIGITTAGG AAAAGGIATG TATATGGICA ATGAAATAAA TACAACTGIA TTTTTAATGA GANTTAACAT ATTTTNNTTT AACAAAAGCA GCATGTAACA CACAATGIAT TATATGT

SEO ID NO:955: (Length of Sequence = 260 Nucleotides)

TATTIGATAG AATTITCIAG IGAAACCATC CIGACTIGGG GITTTATTIT GGAGGAATIT TAAGITATTA ATTCCGICTC
CITAATAGIG ATAGGACIAT TCAGATIACC TTATITCATA TITGGIGAGI TITGGIAGCT IGIGITICIC AAGGAAGIGA
TCCATTICAT CIAAGITGCC AAATTIATGI GIGIATAATA ATTIGIAGIA TICCNGTATT ATCCNITIGA IGICIGIAGG
GICTCIAGIG ATATCCIATG

SEO ID NO:956: (Length of Sequence = 216 Nucleotides)

CCCIATIAAA TCATIAAGCA TIGCATGCAA TACITITINCI GIGAAAATTA TIAACITCCI GGIATATAAA ATTATITICTA GITATGITTA AATATITICCN CIGGGATATI ATCATCITAG ATCIGIAAAG TGGIACTAAA ATAGITAAAA ATTATITINIA AGATATACAC AAACAGAAAA ATATAAAANC AAATGTATCI TATACATAGI ACITGG

SEO ID NO:957: (Length of Sequence = 353 Nucleotides)

TATGIACCAG GIGIGGAGCC TAGAACAGAC ACCAGTCAGA AGIGCAGATA AGGICIGACT TITCCAGCATA GCCAGGGGAC
TITGGCTGACT CCACATGICC CCAGGCCITA CCTAGCTGIA AAGCAGGCAG GITGIGAAGT CATAGIGGCA GITTATGAAA
TATTIAGGGG ACCTAATAAT CITTAAATIG TATAACATIT CITGCATAAA TITCCITICA TGAATCCITT CATGACTTAG
ACCATCIATG ACATGCTIGG ACTITCTGAC TIGICCTAAC CACCCCTCIC TITAAACAAC CAGTCTTTIT ACTITAGGAC
AAGAATTTAC CATACAAGAT TCTTTTGIAT AAA

SEO ID NO:958: (Length of Sequence = 410 Nucleotides)

AAGGAAATGA ATTIGATAGC AGATTGTTAG AGATTAATTA CCTATCATAT GCCAAAGCCA CTTCCTACAT GTCAGTGCTA
AGGAATCCCC TAGAGATGGA ATTCCTAGGT TCAACTGAAA ATTAATTGTA ATTAATATAA TAGGTTAATT CATTGTAATT
ATTTTTAAGC CTTTTGGCAA TGAGTTAATT CCACAAGATC CACATTGCTT GAAGTGTCAC AGAGAACACT TGATGAGAAT
GINCTAGTAA TAAACCTTAA CCCTCTGGGG AAAAAATCCT ACTGTCTTTC CTTCTGGCTT CGTTTCTTCT GGAACATATT
TNGGTGGCAT TTGGATATCT GGAGGACAAA GGGATCCCTA CAAGGTGGNT GCATAAACAT GCGTGGGCCC AGATGGACTG
TGCTCATTGG

SEQ ID NO:959: (Length of Sequence = 197 Nucleotides)

GCCCGGCGAC CGTAGCATCT TCTGGACCAC AAAATAGAAC ATTGCCAGGC AAGGCAGGGC ATTTGGGGAA TTTNAGAGAA AGCAGGATGA GTGATGGAAT TGGGAGGGTG GCACAAGATG TTAAACAGCA TATCTTAGTC CTCATCTAGG GTATAAAACA GGACCCATGG ACTCTAGCAT CCTGGAATGA CAGAGGG

SEO ID NO:960: (Length of Sequence = 345 Nucleocides)

AATAAACTIC TIGITIGITITA AGCCACCIAG TIGITIGICAC TIGITIATIGIC AGCCATTIGGA AACCAACACA COCGCACATG
GOGIGITITAA CECAGECTEA TACAACCITA AGAAAGGAAT GENTGTGGTC ATCAGCAATC TCCAATACCT ACAGCAAATG

GENAGACAGE GAAGGACCAG AGETETAGET AAAGCAAAAA GCCACAGETC ATTAGGAAET GATGCTCCAA CTGGGCATGG AAAAGGAGTT TGGAGTTAGG AACACGACAG ATCTGTCTGG ACAAGGNTCC AGATCTCTCC TAGGGGGAAG NAGGGGCAAC TTAGGACAGT TTTTGTGTCT GTGGG

SEO ID NO:961: (Length of Sequence = 327 Nucleotides)

GCTGAAGAGG AACATGTGTC CTCGGCCACT TCAATCACTG AGTGTGACAA ACTTTCTTCC TTTGCCACAT CAGTGGGTGA

GGACCAATCT NTGGCCTCAC TTACAGCTCC CCAGACAGAG GAGACAGGCA AGAGCTCCCT GCTGCTTGAC ACAGTCACAA

GCATCCCTTC CTCCCGTACT GAAGCTACGC AGGGCTTGGA CTATGTGCCA TCAGCTGGTA CCATCTCACC CACCTCCTCA

CTGGAAGAAG ACAAGGGCTT CAAATCACCA CCCTGTGAGG ACTTCTCTGT GACTTGGGAG TCAGAGAAGA GAGGAGAGAT

CATAGGG

SEO ID NO:963: (Length of Sequence = 278 Nucleotides)

CTCAAACACC CGAGGCCGGG AGGAAAGAGA AGCCGATGCT TCAGAGCAGA CACTCCTTAG ATGGCTCCAA ACTTACAGAG

AAAGTGGAAA CTGCTCAGCC GCTGTGGATA ACGTTAGCAC TGCAAAAGCA AAAGGGGTTT CGGGAGCAGC AGGCGACGCG

GGAGGAGAGA AAGCAAGCCA GAGAGGCCAA ACAGGCAGAA AAGCTCTCCA AAGAAAATTN GAGATCTCCG ACTCGGCTCC

CCCAGCGCCCG CTGGTAAAAG AAGTCACCAA GAGGTTTT

SEO ID NO:964: (Length of Sequence = 349 Nucleotides)

ACACTCTCLE TATAGACAGT CGTGAAGAAC AAGGCTGAGG GATTTINAAG TAAACCCATT TTCAGGATGA CTACAATCCT

TCCACTTCTA GAAAACTTAG AAGTACAAGA AATAGCTCTA CTACGGGTAA CTGATTTAAC AATTTCCCAA ACACCCTTTC

CACTACCCAA GCCCGTGGCC CTCAGAGAGA ACCGCGATGG ATTGCCATCT GGGTTCAGAG GCAATATGAG GAGGTTGGGG

GGATGCCAGG GGCATCCTCA GGGTTGGGGG GCAGGCCAAG GGGATGAGAT GGCAAAGGAC AGCTTTNGGA ATCAGATAGA

CGATCCAGGG TGCCTTCCTA CACTTGCAT

SEO ID NO:965: (Length of Sequence = 361 Nucleotides)

AGCAGCAAGC CAGACGTGAC TGTCAGGAAC AAGCTAAAAT AGCTGTGGAA GCTCAGAATA AGTATGAGAG AGANITGATG

CTGCATGCTG CTGATGTTGA AGCTCTACAA GCTGCGAAGG AGCAGGTTTC AAAAATGGCA TCAGTCCGTC AGCATTTGGA

AGAAACAACA CAGAAAGCAG AATCACAGTT GTTGGAGTGT AAAGCATCTT GGGAGGAAAG AGAGAGAATG TTAAAAGGATG

AAGTTTCCAA ATGTGTATGT CGCTGTGAAG ATCTGGAGAA ACAAAACAGA TTACTTCATG ATCAGATCGA AAAATTAAGT

GACAAGGTCG TTGCCTCTGT GAAGGAAGGT GTACAAGGTC C

SEQ ID NO:966: (Length of Sequence = 163 Nucleotides)

CTGCCTTCTG GGTTCAGCG ATTCTNATGC TTCAGCCTCC CAAGTAGCTG GGATTACAGG CATGTGCCAC CATGCCCAGT
TAATTTTTGT ATTTNAGTG GAGATGGGGT TTCGCCCTGT TGACCAGATT GGTCTTGAAC TCCTGGCCTC AAGTGATCCA
CCT

SEO ID NO:967: (Length of Sequence = 365 Nucleotides)

GTGTCAGTAA TATGTTGTAC ATATTATINC ATCACCCAGG TGTTAAGCCC AGINCCCAAT AGTTACCTTT NCIGCTCCTC
TCCCTCCTC CACCCCCCG CITCAAGTCT ACCCCNGTGT TTTCTTCTTT GTGTTCCTAA GINCTTATCA TTTAGCTCCC
ACTTGTAAGT GAGAACATGC AGTATTTGGT TTTCTTGTTCC TTTGTTAGTT TACTAAGGAT AATAGCCTCC AGCTCCATCC
ATGTTCCCAC AAAAGTCATG ATCTCATTCT TTTTTATGGC TGCATAGTAT TCTGTGGTGT ATATGTACCA CATTTTCTTT
ATCCAATCTG TCATTGATGG GGCATTTAGG GTTGATTCCC TGTCT

SEO ID NO:968: (Length of Sequence = 390 Nucleotides)

GIGTATAGIA ATTIAATAGI AATTAAATGI AGAGIATTIG TAAAAACAAG GAGAGGAAAA AGAACAATTC ATATITGAGA
ACTCCTAATA ATCITCTAGA GCAGAGITCA AAGAAGCAGI GGIAAAAATA AAGCCAAAGA GATATAGGGG CTAGTCTTAG
AACCAGGACT TCCTATAGAA CCAGCTTCCT ATAGAATCTG AACTTTATCT GAAACTCTTT CACAGATCTC CTCCACCTTA
ACTTCCACAA AATAAGAAAT TTGGATTTTG AAGGCAAATT TGTATATTTT AAGGAGCAGG ACAATCTCAG CTGTATCTGG
GTTTGCAGAT ATCCAACAAA TCCTACCCAA ATCACTTTTC CAGCTGCAGA CTTGGAATTT CAGATCCAGG

SEQ ID NO:969: (Length of Sequence = 340 Nucleotides)

CAGACAGAAA AAGATTIGAA GAGACGGGIC AGGAACIAGC GGAATIACIG GAGGAAGAAA AACIAAGIIG TGIGCCAGIN
CICATCITIG CIAATAAGCA GGAITIGCIC ACAGCAGCCC CIGCCICIGA AATIGCAGAA GGACIGAACC TGCATACCAT
CCGCGACCGA GICTGGCAGA TCCAGICIIG CICAGCICIC ACAGGAGAGG GCGITCAGGA TGGCAIGAAC TGGGICIGCA
AAAATGICAA TGCAAAGANG AAATAAAATC TAGACGAATG GAGATGCAGG AGCITCAGGA GCCGAAITCG GGCCITAAAA
ACACIAATIT GCIGCTITCT

SEO ID NO:970: (Length of Sequence = 372 Nucleotides)

TITIAAGATG GGATCICACG GITACCCAGG CIGGAGIGCA GIAGIGCGIC ATAGCICACI GIGGCCICAA ACTCCIGAAC
TCAAACTATC CICCIGCCIC AGCCICCCAA ATAGCIGGGA CIGCAGGCAC ATGCCACCAT GCCIGGCIAA TITITITAATI
ATTITIGIAGA GAIGGGGICI CACITIGIIG CACAGGCIGI TIGCITGATI CITAAGAACG TATAGGGATC CAGCIGIACA
GAGCITTCIG CAGICTITIG TAATAGAATI AGITGITAAA ATIGIACTIA TIACATGAGG CATCAAAGAC CITGGAATAA
AGCIATINCC TCACATATCI GGGCCATTAT TITGGACTIA CIAIGGITAC CG

SEO ID NO: 971: (Length of Sequence = 337 Nucleotides)

GACTATAGAG AACGCTGAAG TITTGAATAA AAGACTCTAG GGTGAGCTTC ATCAGTGCTT GCTTTGGATC CAAGATGTAA
TGAGATTCTN CTTTCACGTC AACAATTGCC GCAAATNCTT TCACCTGAGT GGAGCTCGGA GCACCCAGTC TCTCTGCATA
TAACCAAAAC AAATTTGAAT CCAAAAGGTA GATGTTGAGA GTCTTGTTGG TTCTGCAGCT CAGGCCTGTG AAGTTTGTGC
TAGTCATGTC CACTTCTGGA AAGAGGATAC CTGTNCTCCT CAATGTGAGG GAACGGGAGC TINGGGGCAT CAACCTCACA
TTTTCTTCTC AAGGGGA

SEO ID NO:972: (Length of Sequence = 396 Nucleotides)

TTCCTTTACA TCAAATATCC TCAATGGAAG AGGGGATATT GCACACAAAT ATCATAAAAG CACTACATAT TACTTTCACT
GGAAACTAAT TINCTACATT AGATATGACT GGATAGGATA GAAGTGATGC AGGATTATAA GACATAATAC CATACACAGC

TECAGACTEA CACAAACACC ATTCAGAACA AGAGAGAGGA GTGTGAAGTG CTTCTCAGCT GEGCTCAAGA CCACTTCTTT
CCAGTGCTGG AAAGAGGGGC TGCATGCAGT GTAGGAAAAG CGTGTCTCTG AACTGCCACA GEGTGTTCTC GAAAGGGCAG
CCCGGTCTTG ATGCCACTTC TCCATGGCTC CTGTTTTTGG GGGAGCTCCA AACAAGTGCA GAGAAGCTGC CTATTT

<u>SE ID NO:973:</u> (Length of Sequence = 401 Nucleotides)

TTCTCAAACT TCCAGTTCTC TTCCTGGGCC AAGATCTGGT CCACCACTGC CGTGGCCTCC TTCCCCTGGC GGATGT TCCCCTGGA GCAGAGAAC TTTTCTTCCC AGCAACTTCT TCATCTGATG GGAGGAGGGA ACTGAATAGC TTTCCCCTGGA GCAGAGAAAA ATNCGGACGA GAGTAGCAAG GAATGAGGGG CTTCAGAGAA CTCTNGGATC AGCCCTCCCA CACTCACTGC CCTTTAAGGT ATCTTTGGGG AAAATTAAGG GCTTCTATGA TGAGTCTGGC AGCTNCCCAC ACTGCATTCT CCTCCTGCAT TTTTTTACCA TGCACCAGGG C

SEO ID NO:974: (Length of Sequence = 37 Nucleotides)

TITACAAATG AACCACTGAG CACCTCAGTA CTTAGCTCAT ACCTCATACC TTAGTTCCTT AGTACTTAGC CTTGTGCCAT
CTTGAATGAG ATGGAGTGAA GTGAAGCTCG AAGGAGTGAC AGAGACATAG TCCTTGCTCT CAAGGGGTCT TTAGCCTGGT
CTGGGGGACA AGATTTCCTC ATCTACCTCT TGAAAGGTGG CAGGACAACT CCACACTGGA GTGTTCTCAC CAGCAGATAG
GTGCTGCGGG AGTGTGGCGC CACATTCTTT ATAGCCACAG GCTTTCGTGG GACTTNCCCT GGGGTCCTTC CCTATTTGGC
TGGGTGGACC ATAAGCGGCA AGTGAATGTG GCAAACTTCA ATTCACAATT AA

SEO ID NO:975: (Length of Sequence = 340 Nucleotides)

GACAACAGAA AAAGAAGTGG ACAGCTACCC TAGATTCTAG CTCACACATA ATTCAGCCAG ATAATCATCA TTTAAATAAT
ACCCCTTGAA ATTTTTCAGA CTTTTCACAG CTCTAAAAAC ACAACATCAG ACATAACATC ACACATTGT TCCAAAGGAC
TAAAAAATCAA AAGCAATTGC AAAGTATTGG GAATCACTTT TATGGCTTTC CTAAGGGACA GTCCCCATCT TTCCAAGGAG
TGTTTTTAAA GAAGCACTAA CTCTGGTAGG TTATCAAACT ATTTTT:NAT TCTAAATAAA TAAAAGACTA ACTGAAGGTC
TCAGGTGCAC ::CTTATTTTT

SEO ID NO:975: (Length of Sequence = 343 Nucleotides)

CIGITCCCIA AATATTATTA AAATTITAAA AATTAGACAT TIGGICIAAA TIAGACAGGT AAGATACTAC TGICCITACT
AGATGCTTTA AAGTCATAAA CIGCITCIAT GGCTTTINAT AATTGINCAA CTIGCTIGCT TIAGAGCCAT TGGATTCTAG
GTAAGGCCTA GAGACATTTG GAGTTAGCCA TGICCCCTAG CTATGCTAGA AAGAGTCCGA CATTATCTGT GGITCTGTCC
TGIATCCTAC ACTCTACACC TGATACATAA TIAAAATTAC TTACACTAAA AATAAAAATG GATGCATTTT TTAGGTAGGA
AGGGTATGGG AAATTATAGG TIT

SEO ID NO:977: (Length of Sequence = 265 Nucleotides)

ATCITIGIAA TATCAGIGCC TAGACTAAGC CIGGCGIATA ATAGGCACTC AGAGATITGA AGAATAAATG ACTAAATGAC TGIATCAAAT ACTIGCCCAT TGITIGCIGI TICIGANIIG TACAAGGCCA TCATGATAAT TGATGATCIT AATAATGIGA GAATATGATT CINITACCIT AGIAAGAGG CCATCAGIIT ATIGGATGAT AGITATATGG AAAAAGAAGA AATGCTACTG TGATAAATAT TIATAATTIT AAACA

SEO ID NO:978: (Length of Sequence = 285 Nucleotides)

ATGGTGGGCT GCCCTGGCCG AGGTGGCCAA GATGGCACCT GTTTCCTGCC TCANAAGAAA AGGCACTGAC GCACTGACCC TTTTAGGTTG TNTGGGGTGT GGTCAGTGCC CTCCTGCCTG AGGGTCAAGT CTCTTTTC/A GTCAACTTCA GCACACCTCA

- SEQ ID NO:979: (Length of Sequence = 316 Nucleotides)
- * GIGCGINCAC ACTOTOCTOC TECTOCOCAA ACTOCTCATO ATTGAAGCOG AAGTGGICAA TGAAGGCAGA GGTCATGCGC
 TGCATCIGGA AGTCCATGAA GGCCTGCTGC AGCACAGCCT COTCAGGGAA GITGAACTOC TTGAGCCGGT CGTCCTCATO
- * GICACIGGAG GAGIGIAGGI GGIGGGIGII CACCAGGICC ACCATGITCI TCITGITGGI CICCGCCAGG GGCCCCGATA

 CGAAGGCITC CCACIGCICC TGCIGCICGC TGGGCAGCIC CITCAGCAGC TTGCCGCAGC TGCICIGCAA TTGGGG
 - SEO ID NO:980: (Length of Sequence = 386 Nucleotides)

AAACTGGCTT GCCTTCATCA TCTCTGCAGG GNTCAGTAAA GATTAGAAAT GGATTATTTA CCTTGTTATA CAAATACACC
TCCTCCCTAC ACCCAAGANT TGAGAGGAAG ATGAGCTGTT CCTGTGTTAA CGCCTGANTC AATCCCATTA TCTGCATTTC
TGTGTGTGGT TAGCGCTCCA GCAGCCTAAG GCGGAGCTG GAAATGACAG CCTTGGAGAC GAGGAAGGCT CCAGGGAGGA
CGGAGAGGAA CACCTGCTGA AGAATAAGAC GGGCGCCACC AGCCGGGCTG ATTTTGGGGA ACGGAAGGTA ACAGAGGGTG
ATGCTTCTAA TCGCTTTTAC AAGGTCTTGG AAAGACGGGA TNGCCTTAAC CAACTTGGGG TTTCTT

SEO ID NO:981: (Length of Sequence = 322 Nucleotides)

GITTATIAAT ATTIAAACAT ATTAAAATAA TACATGINCA TAATGAAAAT GAAACATTAC AAATAAATAC ACAGGAAAGG CAGTATICCC CTTCCAGTTC CACTCTTGAA ATAACCAGTT AACAAGATGA TGAACATCTT TCCATGATGT TCTCCAAGAT TCATATTATT TTTGCAATCA TACAATGGCA TATACAGCTC AGGTGCGGTG GCTCACGCAA GIAAATCCCA GCATTTTGGG AGGCTGAGGC GGGTGGTTCA CCTGAGATCA AGAGTTCGAG GCCAGCCTGA CCAACATGAA GAAACCCTGT CTCTTACTAA AA

SEO ID NO:982: (Length of Sequence = 305 Nucleotides)

CCCAAGGCTG TAGTTCAGCA TCAACAGGGC AGGGAGCTTG GCAGGGCAAG GGCAGAGCTG GAGATCATGC CCAGINITCC
AGGTGCCCTC CCTCCCAATC AGCCTGGGGG GCACAGGACA GGGATGGAGA AGGGGCTCTC TCCATGGCTT GGGTAACATG
CCAAAGGCAG GTCATAGGGC AGACTCAGTG GGGGTGGGGG CCTGGCTAAC AAGCAATGGA GAGAACGGGG GCCATCCAGA
GAGGTTGGCA GAAGAGAGCC CCTGGGTCAA GAGAAAACTT TGGGGAAGAC AAGACACGG AGAAG

SEO ID NO:983: (Length of Sequence = 399 Nucleotides)

AGCCCTTGIT TIGITITIAA AAGCIGICGI GITACIGCIT AAAGICTCCA AACTGITATI GAGAACACIG ACCAGAGCCC
TGICCATAGA CCAGIGITIT TCCAAGIGCA GATIGCAACT CCITIGCAGA GIAGGITGIG GAGCCATTIN AGCIGACIAC
TCACCAGCIT TCTTCAAAAAT GIAAATGGAA TAGGATAGAA AAATAATGAA AAATTGIAAA GIGAATTGGA TGCAAAAAGG
GIAAATATIG TNGIGICAGA CITITITIGGG TGAGIGIGCA TGIGITCACA TACTGGNTCA CATTATAACA TGIATTGCTC
ATTATGGGT GIGGICAGAA AAAATTCAGN AAACGCTGIC TCAGACTGIC CCCAAGITGI ATTIGCTTAT AATGGGACT

SEO ID NO:984: (Length of Sequence = 408 Nucleotides)

GIGGIAIGAG GIATCAATGA AATACATITA AGATGIACAT TGGITTGITT CAGAAAGGCG AGACAAGICA AAGCGGGGAC TICCAGGCIA TAGGIAAAIT TATACATITC CIGGITAAGA TIGGITGAGI TIGICTAAGG ACCIGGGATC AACAGAGAGG AAATGITTGG NITAAGACAA GGATTGIGGA GACCAAAGIT TIACTACGCA GAGGAXOCIC PTAGCTAGCA GGCATAAGAC AGAAGAGGCT GIAAAATGIT TICTTATGAG ACTGAAAAGG GIGCCTGACT CITAATTGAT TATCTCCTGG NICTGGAAAG AAAAAAAAAA GGGAATGGCC AGGTGCGGTG GCTCAGGACG GGTCTGGTGG CTCACACCTG TAATCTTTCT TAAAACGTTA
TGAAGTTC

SEO ID NO:985: (Length of Sequence = 439 Nucleotides)

TOGIATACTI TIGINITITI TICIACTIGI TAGIIGIATI AGIACAAAT GGCATAATAA AGITACTIIG TITIGCCATIT
CCCACTCATC TGAAAATCAC AAAAAGCAIT TATTICTAAG ATITATATCC ACTGACCTIT TCCCCAAAGI TATTITCCIG
TTACTIGIAT TICATCTIIG CCCTTATITC TITAATATTI GIATTAGAAT TAGCTIGCTC TIGITICCTI CACGGCAAAT
GIGITACATT GCCCACTGGG TGGCTICTGC GGATGCCCCT ACCCACCCCT CGICTGGAGC AGAGAAGICC TGITAGCCTA
GCAGCATAGI GGCTGCTGIC AGIGCGAGGA GITGIGCTIC TCTAGCATGG TCTGIGATGI CATCTGGACA TAATTAATTA
GACTAATCCG AATAGAGGAC CAAGACAGCC CTGCCTGCG

SEO ID NO:986: (Length of Sequence = 286 Nucleotides)

COGCGACGAA CATGGAGAI CCAGCTTG GAGCGCAAGT CCTCCTGCGG GAAGAAGTGT CGCGGCTCCA GGAGGAAGTT
CACCTTCTCC GGCAGATC "AGATGTTG GCGAAGGACC TGGAGGAGTC GCAGGGCGGC AAGTCCTCTN AGGTCCTCTC
GGCCCACCGAG CTCAGGGT "TGGCCCA GAAGGAGCAG GAGCTAGCCA GAGCCAAAGA AGCCTTNCAG GCCATGAAAG

CTGATCGGAA GCGCTTAI CGAGAAGA CAGACCTGGT GAGCCA

SEO ID NO:987: (Le of Sequence = 381 Nucleotides)

TCCAAAGGTT TTCATCTC ::TGGATAA ACAAAC TG GTACATCTAC ACAATGGAAT :TTGGGA GATGAAACAG

AATGTATGAG GGGCCAC: ::CATGTAT GGTE ::XG GTCTGCCTCC CA: TNTCCA ::CAGGCA G: TGTGCT

GGGTGAGGGG CTGGGAGG: ::CAGGAGG CATC: :::AC AAGGGTGGAA GC ::GAAGA ::CGACCAG TL.:ACAGGGT

GTNTCACATG GTACAACCAA GAGACTTGGC GTGGTTGGG GGGAGACTTG GAGGCCCCCTC T

SEO ID NO:988: (Length of Sequence = 381 Nucleotides)

GAATTAATAC CAATAGAAGG GCAATGCTTT TAGATTAAAA TGAAGGTGAC TTAAACAGCT TAAAGTTTAG TTTAAAAGTT
GTAGGTGATT AAAATAATTT GAAGGCGATC TTTTAAAAAG AGATTAANCC GAAGTGANIT AAAAGACCTT GAAATCCATG
ACGCAGGGAG AATTGCGTCA TTTAAAGCCT AGTTAACGCA TTTNCTAAAC GCAGACGAAA ATGGAAAGAT TAATTGGGAG
TGGTAGGATG AAACAATTTG GAGAAGATAG AAGTTTGAAG TGGAAAACTG GAAGACAGAA GTACGGGANG GCCTCCTTCA
TGTTTACAAT TTTAATTAAT TTTTTTTTATT TTAGGNGTAA TTTCTTACCA AACATTACCC A

SEO ID NO:989: (Length of Sequence = 432 Nucleotides)

GICTIGGUE ATGCCICCI CIGCUTCCTG GGITCAAGCG ATTCCCCTGC CITAC ACC CAAGIAGCTA AGACT 3
CATGCGCOO ATGCCTGGC TAATATATAT ATATATTITI NGTAGITTIA GIAGALACGG GGITTCACCA CGTL 3
GCTGGTCTCG AACTCCAGAC CICAAATGAT CIGCCCGCCT TGGCTTCCCA AAGIGCTGGG ATTACAGGCA TTAGCCACTG
TGCCTGGCCA ACAATATATA TTAAATAAGC ACACATACAA CAAAAGIAGG TGTTGGTAAG CITACAAAAA TGTGACCAGT
AGCTTGCTGA AACCTAACTT TTTATTTGTT CATGGAACTT TCTAGACCGT AACTACACTG AATAATGAGA ATCTGCTGTA
ATCTTTTTTA GGTGCTGTAG ATGAGCCATT GG

SEO ID NO:990: (Length of Sequence = 421 Mucleotides)

GGCAGCCCTA CTTTINCTTC TCATTAGCAG TTTCAGTCCA CAGCTGGGGT ATTAAATTTG TNAGTCATTG AAATTAATCC CTGACTGAAT TGGAAAGGAA TTGTATTTGC AGTATTTGGA 1TTATTTATT TINCAGGTAT GGAATTCTGG TGATTTTGAA AACATGAATG ATACCATTIT GCAGCAGCAT TGTAGATTITG TAGTATITTA GATTGGTATC ACAGTGCACC TGAAAAGTAA
GTTTCATTIT ACTITITINA TIGTIGITGA GACGGAGCTC ACTITIGICA CCCAGGCTGG AGTGCAGTGG TGTGATCTIG
GCTCATGGCA GCCTCTGCCT CGCTGGGTTC AAGCGATTCT CCTGCCTCAG CCTCCCGAGT AGCTAGGACT ATAGATGCTC
GCCACCATGC CCAGCTAATT T

SEO ID NO:991: (Length of Sequence = 351 Nucleotides)

CCTCACTCCC CGCGCTGGCA CCTCAGGTT ACAAGAAGAA CTAGGAAATA ATGCCGGCCA CGCGACCCCT GGAGAGGGGG CCGGCTAGAA CAGCGTTCCT AAGAATCCGC GCCACAGCAG GTCCCGCGAT GTTGGGGCCT TAGTGTCATC GAGCTAGCCC CAATCCTCAA CCCGATCTTC AACTTCTGGT AGTCCTAACA GAAGTCTCGT ATTGAACCAG CCACTINTGGC CAGGGAGAAG TAATCCTCTG ATAGTTGAGG TTCTTTNCTC TCCTCTGGAG CAGATAGTGG TGTCTCCTCC CCACAAAGCT CATGTTCTGC TGGAAGAAAT GGAGATGGCG CCCTGGAAGG C

SEO ID NO:992: (Length of Sequence = 406 Nucleotides)

CCAGAAAAA TGGCCACTAC TACCACTIGG CTCAGAAATG CTAGTCTTTA TITINCTGAAA TGTTTTATAT AGAAAAAATT
TAATAATAAA TAGACATTCT TATATATTTC CTTACCATTT NAGATTGGGT TAAAAAGTAT GGNGACTTCC GGCCGGGTGC
GGTGATTCAA GCCTGCAATC CCAGCACTTT GGGAGGCCGA GGCAGACAGA TCATGAGGTC GGGATCTGTG GCTAACACAG
TGAAAACCCCG TCTCTATTAA AANTACAAAA GGAATTCCTG CAGCCCGGGG GATCCACTAG TTCTAGAGCG GCCGCCACCG
CGGGGGGGCT CCAGCTTTTG TTCCCTTTAA GTGAGGGGT AATTTCGAGC TTGGCGTAAA TCATGGTCAT AGCTGTTTCC
CGTGTG

SEO ID NO:993: (Length of Sequence = 381 Nucleotides)

ATGGAAGGAC CGTGCCGGGA CCCCAACGAG GCANTGCGGG AGITTGCCAA GGAAATTGAC ATCTCCTGTG TCAAAATTGA
GCAGGTGATC GGAGCAGGGG AGITTNGCGA GGTCTGCAGT GGCCACCTGA AGCTGCCAGG CAAGAGAGAG ATCTTTNTGG
CCATCAAGAC GCTCAAGTCG GGCTACACGG AGAAGCAGCG CCGGGACTTC CTGAGCGAAG CTCCATCATG GGCCAGTTCG
ACCATCCCAA CGTCATCCAC CTGGAGGGTG TCGTGACCAA GAGCACACCT GINATGATCA TCACCGAGTT CATTGAGAAT
GGCTNCCTGG GACTCCCTTT CTTCCGGCAA AACGATGGGC AGITTCACAG TTCATCCAGC T

SEO ID NO:994: (Length of Sequence = 384 Nucleotides)

GITCTICCAG TICGGAAGGA TAAAATCAAA TICCCACTIT CIGGGGIGGA TGCCCAAAAC CITCACAACI CAAGIGITCI CCAAGIGCAA AIGICAAAAT GGGAGGAGGA AAGGGITTAA AAATTAGAGA AAACTGIAIG CACITACGGA CITAAAAATC CGAAAAAAACAI AGIAAAAAACA TAGCAITAIG CICTGAAAATC ACAACCAAAG CCAAAATAAA AGGGACAITIT TICCACCTAAA CIACCIAGAG GGATTITITG TITAGITTIT CCITTITCIT TITTITICA TITTCCAGIT AAGICCTAIG TCITINGIGA AATTCCAATA CITAAACIGC AAGICGCAA TCGICCICGA AGICAGIGAA ATTA

SEQ ID NO:995: (Length of Sequence = 386 Nucleotides)

ATAACTITAA CAGAGGATTG GAATAATGAG GGATTGGCAA GGAAGCAGTA AAAGGGAACA CTAAAGTATA GAATAATAGC
AAACAGAAGG AGCACCCTAC CCCTAGGGCT GAGAAAGAGC ACAGGGAAGT CCTTTTTINT TCCTGGACAG AGATCCAGAC
GAGCTGGAGA AAGAAGTTGC TATGGTACTG CATCANTGGA ACTTGCTGGA AATCCACCCT CAAGGGCACT AGGAAAACCT
GTTCAGGGGA GCTGTGGAGG GAAATGGGT TGGCAGGAAA GCTGCTGGGC GCGGGTGCT TCAGACTGCA GTGTATTGCA
GGAGCTTGGG CACTGGGGAA GCTGTGTGCA CTGCAGGATC CTGCTGAGCC AGCACATCAG ATCAGG

SEO ID NO:996: (Length of Sequence = 307 Nucleotides)

GIGCGCCAAC TGCAAGAAGG AGGCCATCIT TTACTGCTGT TGGAACACCA GCTACTGTNA CTACCCCTGC CAGCAAGCCC ACTGGCCTGA GCACATGAAG TCCTGCACCC AGTCAGCTAC TGCTCCTCAG CAGGAAGCGG ATGCTGAGGT GAACACAGAA ACACTAAATA AGTCCTCCCA GGGAGGCTCC TCGAGCACCAC AATCAGCACC TTCAGAAACG GCCAGCGCCT CCAAAGAGAA GGAGACGTCA GCTGAGAAAA GCAAGGAGAG TGGCTCGACC CTTGACCTTT CTGGCTCCAG AGAGACG

SEO ID NO:997: (Length of Sequence = 402 Nucleotides)

TCTGCACCTA ATACTGAGGG TGTGAAATCT TCCTCAGTAA TGCCCAGCCC TAGTACCACA TTAGCGCGGC AAGGCAGTCT
GGAGTCACCG TCGTCCGGTA CGNGCAGCAT GGGCAGTGCT GGTGGGCTAA GCGGCANAGC AGCCCTCTCT TCAATAAACC
CTCAGACTTA ACTACAGATG TTATAAGCTT AAGTCACTCG TTGGCCTCCA GCCCAGCATC GGTTCACTCT TTCACATCAG
GTGGTCTCGT GTGGGCTGCC AATATGAGCA GTTCCTCTGC AGGCAGCAAG GATACTCCGA GCTACCAGTC CATGACTAGC
CTCCACACGA GCTTCTGAGT CCATTGACCT CCCCCTCAGC CATCATGGCT CCTTTGTNTT GGACTGACCA CAGGCACTCA
CG

SEO ID NO:998: (Length of Sequence = 304 Nucleotides)

GCAGGETTET GATTETNAAG ACTCACAACC ATGTGGAGAG GCCGAATCAC GCAGGAGAGC CACGCATTGG AGTACCCTGG CTCCCAGCCC CTTCCCCACC CCGINITGAG CCAGAGAGCT ACAAGCAGGA ATCCCAGTGC AGCTGCAAAT NATGGCCATC GAGGAAGTCT GTGGAGAAGA GGCTGGGGC TGTGGTGCTG AGGGGGGCTA GGCTCAGCAC GGGACCACCT GACGACAGCT CCCAGCCAGT CCATGCTGTC CAGGTGGCCA TCAAGCCAGG TTCCAGGGCC CATGGGTGCT TGCT

SEO ID NO:999: (Length of Sequence = 321 Nucleotides)

AGAATGGTTT TGGAGCTCGA NATCTTCATG GGTTAGACTT GCTGGTCAGA CCCAGGAGCA CCTGTGGCTC ACACCTTCTG
TNCCCCTCCT GGCCTGTGCA GAATGTAAAC AGCAGACTCA TACTCAATGG GCACTACAGG CCTTATCAGA CGTTTTATAC
AAGCCTGGAT TGCTTAGTAG GGGAATAAGG CATTCTCTGA GGGGGCTTTC CACTTAGATT GAGAATTTTA TTTGAAAAGA
ATCTGGTTTA AATGGCATTG TGGTCCGAGG TAGCTGCTCT CCCCACTGAG AGCTGAGCCG AAATATAAGA ATAATATATT

SEO ID NO:1000: (Length of Sequence = 253 Nucleotides)

CCCTAGAGGA TITIGCCGTCT TINATCIGCC AGTGACCTGA ACCACGCAGA TITITICAAGC AGGAGGGCCG ATTGGGCAAC
CACAGCTCCC GIGCICTCTC TITIGCAGTIGC GCGGCTTTCC CTCCGAGAAG GACTTTGAGG ACTACATTAG GIACGACAAC
TGCTCGTCCCA GCGTGCTGGC CGCCGTGGTC TTCGAGCACC CCTTCAACCA CAGCAAGGAG CCCCTGCCCGN TGGCGGTGAG
ACGTGCGGCC GGG

SEO ID NO:1001: (Length of Sequence = 164 Nucleotides)

AAACAGAGTA CTGGGATGTC ACTGTTGGAA AGTGCTCACA ATTTCTCATC TAAGCCGAAG TTGTCTGTNC TCCTTCCTAC CTTAACAGTT TCTCACTGCC TGAAGGCAGC TGCCAAAACC CCTCTAAGCA AGCAGCACTC TTACCCACCA AAATCTATGA CCTC

SEO ID NO:1002: (Length of Sequence = 262 Nucleotides)

ATATCITCCT GAGGGAAAGT GGTAGAGTTA AAGAGGGCAT AGAGAGCGCA CICATGCATT TACAACTCAG AATITTAAAA AAAGITTACA TITTGICAIT TGTACTTCAG ATGAATTINC TIATTAAAAG AAATAAGGCC ACAGAGGTAA ACTTAAGICT CCTGTTTCCC AATGCCTACC CTCCTTCTTC TCCTTTCCTC TTTCTCTTTC CTAGAGAAAT CCTGCCTTCC TTTCCCTTCC CAGAGGCAAC TGGCATTATA AT

- SEO ID NO:1003: (Length of Sequence = 267 Nucleotides)
- * GGAAAGAGA GCAGGICIGG AGGITTGIGG AACCCAGTCC CCIGCAGAAI CIGIAAAACC TAATAAATCA TGGITGIGGC CATTCICACG GIGGIGAITG TAATTAGACG ACCCCGGGA AGCCCAGACA CICGGGGCCT GGAGITCCIC CCCCIGCCIG
- + ACCTAGAAGC AGAACCGITT TCAGCGVICT GCCCTGITGG CTTTAAGGCT TTGTCTTAAT TTAAGGAAAA AGATCCTCCC
 GGGTTTTATT TCTCTCTTC TTGAGTG

SEQ ID NO:1004: (Length of Sequence = 277 Nucleotides)

GGCTCCTAAA CACITTCTTC CTGAGATGIT AAGCAAAGTA ATCATCCTGT CACTAGATAG AAGCGATGAA GATAAAGAAA AAGCAAGTNC TTTGATCAGT TTACTCAAAC AGGAAGGGAT AGCCACAAGT GACAACTTCA TGCAGGCTTT CCTGAATGIN TTGGACCAGT GTCCCAAACT GGAGGTTGAC ATCCCTTTGG TGAAATCCTA TTTNGCACAG TTTGCAGCTC GTGCCATCAT TTCAGAGCTN GGTGAGCATT TCAGAACTAG CTCAACC

SEO ID NO:1005: (Length of Sequence = 271 Nucleotides).

GTTAGGTCAT TCACACATGG TGGAGACAGG AATCTACAGA CTAGGGATCA GCCCCAAGGC TATGATCTTT GINCTGCGCC
GCTCTACCCC TGAGCAGACG GGCCAGAGGT CCAGAGAGGG CTGTGCTGGC AGAGTTCATA CTTTGATAAC TGAACCCTAG
AGTAAGCCTG CCCTGGGAAA TNCCAGCTCA AGGGACTGAC AGGCATAATG CTCTTTGGGA GAGAAATGCC ACATCTGCAG
CGACACGNAT CCTTAACACT GTCCCAGGAC T

SEO ID NO: 1006: (Length of Sequence = 336 Nucleotides)

TATTITICAG ATATGGATAA AAATTGCTTA GGAGAGTAAA GAGAGACAAA GITGAAAGCA GGITTATAGT AGGIGITGIT
TTAGTGTTGA TCCCTTTTTG CTCCAATAAT CAAAGTGATA AATATGAAA ATTGATTCAT GCAGCATTAC TTACTCCATT
CTAATTTTNA TATATGTCAA AAGTGCCATC TCCCAAACTG TGCTATCCCC TTCAGGAGAA GAGACTCTGC TGAAGTTTAT
AAGGTTGACA TATTGCCAGC TTCAATAATG TAAAGATGAA GTGTATACTG GAATTCTTAA TGCAAATAAC AACTCTTTTG
GGAAGTAACC CCGTTT

SEO ID NO:1007: (Length of Sequence = 355 Nucleotides)

GECAAGAAG CETCEGCEC GCANTECEGA TOCAGAAGGA CATAAACGGC AGCTTGITCC TOCAGGCTGG TEGGCTINGT
GCCCTCGGCC TTGGGATGCT TATCACAGIC CTTTGGGACC AGAACACTGG ATATCAGINC AGCCTCTGGG CCAGCTTCAG
AGGCTGITAG AGCATCATTG CTGCTGTGGC TGATGCTTCC TTTCCTCAGT AAATCACAAA AGTCGTGTTG GCCATCCAGG
TTACCGAGTG ACTTAATTTC CAGAAAATTT AATATTGAGG TCATTATTGT ATGCATTTTC ACTGTTGCCA TTTTTGTATC
CTCGTAGGTA GGTCTATGAA GTACCACTGG GGTCA

SEO ID NO:1008: (Length of Sequence = 269 Nucleotides)

ATATITAAAG AGAGCTITGG TCAGIAAAAG TATAAAANCT GAGCTITGGI AAGGGIACAG TITATAAAGGC CIAGAGAACA

**
TCAAAACATI CATTICATAT TGAATGIATA AATACCCACA TGTGAGAGCA CATGTTGATT CAGITTGAGT ATGTCTGCCT
TGTGGNICIT TAAAACCTIT CCAGCCTGGG TTATTTTCCC AAGCTTTCTT TATAATTACA CCAGGGAAAG AGTTACCNGG
NATTAATCAA AACCAGACAG TGGACAATG

SEO ID NO:1009: (Length of Sequence = 295 Nucleotides)

GATAGCAGCA ACATACGITT GITTATICAT TIGCITACIT ACAACAAACG TITATICATI ATITATAATG CAACAAGCAT TAACCTAGGI GCTAAGGAGA GAAAAATGAG TAAGACACAG TITCITTCCT CAAGGAAATC ACAGTCTGIT GGCAGAGATA AGITAAGTAATG GTGCCTAATA TAGGTAACAC TIGCTACCTG CTCCAAGGAAC AAAGTTAAGC AAGTGATTAA GITAAGCAAT GCTTAGAGGI AGAGGATGIA AGANTGGCCT TAAAAAATGT GTCTTCTGAG ATGAG

SEO ID NO:1010: (Length of Sequence = 356 Nucleotides)

GIATTICCIC ATTIGIGCAA ATNAAATAGA AAAGGTAAAT NAGAAACICA AGAGGTIIGI TACCIACIGI CAATGGAGIG GGGAAAATGG GIGGAAAGAA GAAGGCAATA AGAAAAGAGI AACAGGAAAC GACAGINGAC ACITCIGAGI ATACCITGIG GAATCICITI CACICITAGA ATCATAGIAA TAGANGANGA AAAAAGAACI CCCCAAACTG AAAAGGATAG ACCACIGGAA CAACITCAAG TGGICIAATG TAGAAGCAAA TGGAGICCCI CAAGGAAAGA AGAGAGGITT TGAAAAGAAA AAAACATTIG AAGAGITAAC AGCGAAACAC TITCCAAACT TAAAGG

SEO ID NO:1011: (Length of Sequence = 315 Nucleotides)

AGAGAGACAC AACTGTAATA GAGACACAGA GGAGTGGCAC ACAGAGACCA CCTCCCAGCT GGAGACAGTC AGGAAGGACT GAGGGAGAGG GGACAGCCAG GGCTCCACAC CCAGGCAAGA ATGGGGGAGG GCCTGTGGAA CAGAGAAGTC ATCAACACAC ACAGTTCAAA GTCTACCCTA GGCTAGGAGG GGGAGCAGGA AGAAGGGGCA GGGACGCAGG GGCCCGGCCT GCNAGCTCCC TGTTGGCCTC TNCTGCCCCC TGCTGGCTCC CNCTGCGGTG CTCAGGCAGG AAGAGAGGAG GCTGCTGTTT TTAGG

SEO ID NO:1012: (Length of Sequence = 272 Nucleotides)

COCAACTOTA TAGOCCIAGI CAACCACTAA TOTATACCOT GINCICIATA GATTIGOCTA GICTAGAAAT TITIGTATAAA TGAAATGCAT GCACTIGAAC TITITIGIATO TGGCTIGCIT TICCATTTAG CATAAAGTIT TAAAGGTCON CATATGTIGC TGCATGIGIG CATTICITIT TGIGNACIGC NATATTACAT TGIATGGGAT ATACCATTIT GCCATATTIN GITAAATCCA TICATCCAGI TGGIGGGACA GCAGGITATI TC

SEO ID NO:1013: (Length of Sequence = 252 Nucleotides)

TITGITAGIG TITTCTACAC TACACICAAG TICATICAGC ATGICATITC AACAACATGI GACGIGICAA CITCAAAAAT TAAACAAACC AGCNAAACAC AACACITGIC ACTACAAAGG AACITGITIT ATTCICAACC TICIATGATA GCTAAACITC TCIGNAATIT NGITCCCCCA CACATCCCAC ATCIGGGCTC AATTTCCAGC TICIGITNIT CIGITTITATT TCATCCAAAA TGITATTITA AT

SEO ID NO:1014: (Length of Sequence = 210 Nucleotides)

GGGATACACT GACAGIAATG TGAAGCGCCA CACTTGCAGA TTTCAGGCCC AGCAGGTCCT GGNCAAGTGC CATTCCACCC
GGAACTTTTA ACCCAAGCGG TGGGGAAGGA AAGCCAAAAC TCCAAGCTGG CACTTTTTTG GGGTTCTGGG CCATGACACT
TCTTAGGCCT TCTGCTGCTG AACTTTTACA GGGACAAAAG GTACCCCACG

SEO ID NO:1015: (Length of Sequence = 222 Nucleotides)

GENAAGAAAG GITTCTCAGA GGACAGCCTT ATTAATTICT CAGAGGATGA ATTTGNACAA TGGCAGCACG TTGCAGTCAC
AACTTCTTAA GGTGCTTCAG AGGCTGATTG TTCCTAGNAA CACAGAGTAA TGAACTATTC CTGAAGAGCA ATGAAACAGG
TTTTGAATTT TNTTGTATCT GNACTTAGNA ACACATCAGT CCCCATCAAC CCATGGACTT CT

SEC (T) NO:1016: (Length of Sequence = 236 Mucleutides)

GAATAAACTG GTTTGGAACC AGAAAAGTAC AAAAAAGAAC AGCTAGAGGT ACATAGACAC AGGACAATTA ATCAATTTGG GAAAAAAAAA AGNACTTACT TTCTCCATTG CTGCCTGAAT TGTTTCCCAA TCTGCCTTGA AATGCCACTT TTGGCCAATA TTTTTNCAAA AATTTGACCA AAAAAGAAAA AGCACTNAAT TTCCCTTTTT ATACAAAAAT GNITAAGTAG GCAAGT

<u>SEO ID NO:1017:</u> (Length of Sequence = 259 Nucleotides)

GCTTCCCTAG ATTTTCCCT AATTTTGGAC CTATGIGGAC AAAAAAAAAA ATCTAGICCA AGCTTCACT ACCTTCTTT
TTTATTCGCC TTCTGCTTCT GNGTTCCACA TGGGAACTTG AAGTGGTTTA TAAGAATGCC ATGCTGTGCA AATAGTAAAA
ATGAATTTNC TGATTTTTAA AAAAGCCCTC AGGAACGGCA TATGTATANG GTATGTATAT GAAAAAANGT GTTNAGGAAT
GCAGGAGGGA AACTAGGCG

SEO ID NO:1018: (Length of Sequence = 354 Nucleotides)

CTGGAGGAGG AGAAGAAGCA TCTGGAGTTT ATGAATCAGC TAAAAAAATA TGATGACGAC ATTTCCCCAT CCGAGGACAA
AGACACTGAT TCTACCAAAG AGCCTCTGGA TGACCTTTTC CCCAATGATG AAGACGACCC AGGGCAAGGA ATCCAGCAGC
AGCACAGCAG TGCAGCCGCG GCTINCCCAGC AGGGCGCTTA CGAGATCCCC GCGCGGCTGC GGACGCTCCA CAACCTGGTG
ATCCAGTACG NCTCGCAGGG GCGCTACGAG GTAGCTGTGC CCCTNTNCAA GCAGGCCCTG GAGGACCTGG AGAAGACTTC
AGGACACGAC CCACCCGGAC GTGGCCACCA TGCT

SEO ID NO:1019: (Length of Sequence = 393 Nucleotides)

GATGACCGAT TIGGCCATGG AAGACITATC TICATGGCAC AGAGAGNYTG TSCAGAGATG AGICAGACIC AGGGGCTGAG
TAACAGCAGA GCAGAGAGTG CAGAAGTGGA CGCTCAGAAG CGAGTTTATG TGTGTYTTTY CCTCTATCTG CTGGCTGTGG
CTGGTACTGC AACCTATCCC AAAGTAACAG CCTAGTCAAT GAGGTATATG CTTCAGATCT GGCAAACTCT CTCTGCACAT
AAAACTGTTA TICTTAGTTC TCTGAAAGAC CCCCACATCT TTGAAGTGTA AACTAAGAGC TACATTTTCC CTTTTACTAC
ATCTCCCTTA AAAGGAAAGC ACTACAAGAG CTTTAAAATA GCAAGCTTCC CTATTCTAAG GGGAAANAGT CTT

SEO ID NO:1020: (Length of Sequence = 403 Nucleotides)

CTGAGGAAGA GAGGIGAAGI GGCATCIACC CAAAACACCI GIGIACIGGI TAATAAGGIC GGIAGITCCC ATTAATGAGC
TTGATGAAGG ATGGCACCIG ACAGGGCCIT AAATGANCIG ATGGAGIGAA TGINACCAGI GIGAATIAAA TTINCITIAT
ATATAATAAA TAGCIGIGCI TACACATITI CAGATITNCI TTGICAGCIA TGGACATGGA ACAGCGGGAC TATGATICIA
GAACCAGCACI CCATGIAGCI GCIGCAGAGG GIGAGACCGG AACIACICCI ATCIATITCC TTTCCAGATI TAATITICIAC
TTAGIACIAA AATCIGCICI TTTTTTGGGG GIGGGACGGI ATAGGICATG TTGAAGITGI TAAATTITIT NCIGGAAGCC
TGC

SEO ID NO:1021: (Length of Sequence = 452 Nucleotides)

ATCGCAACCT GECAGGGGIG TGGGGTTTGC TGGGGGCCTC TGTGGGGCCA TGATCTGAGG AGGGTATGTG GGGGGGGGGA GCTCAGCACA TTCCATGGCC TAGAGGGGCC ACACAGAGGC CCCAGTGGGA CCCATGGCGT GGAGGCAGGT ATGGGGAGTT KTGGGGAGAT CCCAGGGTGG TCTGGGGCCT GGAACCGGCC ATTKGGAGGC CCCAGCAGTT TCAKTGCCCA GGGCCTCCCT GCAGAGCCAT GCATGGCAGA AGAAGTGTGT AGCATGAGCT GGTACACGCC CATGCCCATC AAGAAAGGCA GTGTGGTCAT GCGTKTGGAC ATCAGCAGCA ATGGCCTGGG GACCTTCATT CCAGATAAAA GGTTCCAGAT GATATCAACG GCTTCCTGAA GAGAGACCCG GGCAATAACA TCCATTCANT TGGGAGAGGA GGTCCCCATT NT

SEO ID NU:1022: (Length of Sequence = 413 Nucleutides)

AGCAACAGAA GAAAGGGCCA CATATATGCA AATGCCTGGT CACTATATCT GGCCCTGAAG AAGGAAGGAG TTTGCAGGGC
TCAGGAGACT GGAAATTTT NCCAGGAGCT AGGAACGAGG GGTTGGGAGA CGTTGGTCAA AGGGTACAAA GTCCCAGTTA
TGCAGGATGA ATAAGTTCTG AAGACCTAAC ATACAGCCCA GTGACCATAG TGAATAACAC TGAATGANCA GTATACTCGA
AATTTGCTAA CAGAAGAGAT CTTAAGTGTT CTCATAACAC ACAAAACATA GCAACTGTAT GAGGTGATGG GTATATTAAT
TAGCCTGACT GTGGTTATAC ATTTTATCAA AATGTCACAC TGTGGCTGAG TNCAGAGGCT CATACCTATA ATCCCCANCA
TTTTTGGGGA GCT

SEO ID NO:1023: (Length of Sequence = 379 Nucleotides)

TCAAGICICA AAACITIAAA AGACAGIAGA TATTIGIGGI TITCIAGCIA AATGAGGGCC AAGATIGGIC TITTICAACI AAATTGAATC ATGIAGIATA TCTGATTICA TAGCITICIG GGGGAAAAGG GAGGATITGA ATTAGCAGCA GTGCAGGICA GGAGCAGIAA AGAAGACAGI AGGAGGAGIC CAACIACAGA TGIGAATGAN CAGCCICAGA GGAACACATG AGAAGGIGAC CTGCTGITTA TCAGGAAGGC GGGGCTTTCT CTCTAAGATA CAAACCAAAT AGGAATCGIC AAATAGITCA AATTATCCGG GGGAAAAAAGC CTGAGCAATG ATCCCTCTGG AAAACAAAGC AGITCTCAGG CAGCACCTT

SEO ID NO:1024: (Length of Sequence = 320 Nucleotides)

AGICTACAGG AACAAAGAA TCTAAGATGG CTGCTCAGCC TTGAAATGTA CATGITTTGC AGCAAAGTTG TTGAAGAACC
TTCCGTTGGC ACAGATTGTC CTTTTTCACA AGCATACAGA AGCCTCCTTC CGCCCAGGNC TCTTCCGTTG CATCCTTGCA
AATGGCTCCC ATTTGACACA TTCCTAAGTC TAAGAGATAC CCACTAGGGC AGCTTGTACA GTTCTTGAAT CCTGGGCCAT
TGCACGTCAA ACAACTGATA TCACATTTTT TTGCAGGACT TGTATCCATT CTCTGAAGAG TGGTCAAAGT AATAGCTGAT

SEO ID NO:1025: (Length of Sequence = 368 Nucleotides)

TATITAATCA TICTITICIT TGCCIGAAGA CITAAAACTA AGAAGATTAT TCGAATGGIG AATTAACTIG TIGAAGAGAC TATICCAAAG GGATAGAATG AGACTAATTY CIGACTATGI TITGCIAGIG ATGGGIGGAT GGGAACAAAC ATTACAAGAA ATAGCATAAT GAATGTAGAA AATATITCAG TITGGAGATG TGCATGANIT AGITTCCTAG GITTGCCACA ACAAAGCATC CCAAACTGGI GGCTTAAAAA ACAGAAATTI GITTCATGGI TCTTGAGCCT AGAAGGTCAA AATCAAGGIG TIGGCAGGAC CATGCCTCTC CTGAAACTCT AAGGGAGAAAC CGTTCTTTGI TTCTNCCT

SEO ID NO:1026: (Length of Sequence = 379 Nucleotides)

GERGEAGGER CATACAGGAA GGACCATGER GECTCAGAGC AAGGGGGGG CCATCTCCTA GCCAAGGAGG GAGGGCTCCA GGGACCCCAA TCCIGCIGGC ACCTAGGCCT TGANCTICCA GCCTCCAGAC TGGGAGAAAA TAACGICTCA TIGITAAAGC CCCCAGCAAA TGANIACAGA ACCTAGGAAG GGGCAATGAA TGANIGATAG GTGGAAGGGC TAAGAAGAAA AGAGGAGGGA GAGGAAAGAG ACGIGCTCAG ATCTGTCTCT NCTGGACATC CGATCCCAGG CTGTCTCTTC AGTGGGNCCA AGTCCAACTA GCAGTCAGCT CAGAAATAAT CCCTNAGGCA TCGAAGCTTT CACAAAGGAG GNCACAAAA

SEO ID NO:1027: (Length of Sequence = 411 Nucleotides)

GCCCTTGGCA CCTAGAAGCA GCCAGGAGG AAGTACTGAC CATTTAAAAG TGGCAGATCT CCGGGCCCCA TTTCTGCAGC CTTCATTCTG CAACTCCAGG GAGGGTATTT TINATTTGTG GGTTCAAAAA ATCTGTATAT ACAGTCTATG TGTTTAGAAT TTGTGTTGTA AGTAAAACTAC AGCTTTGAGT TGGAAAGAAG TCACGGGTTG TAAAACCATT TGGATTTTTT TAAAACAAAA GTATTAATAA TCTGGAAGAC AGTNTGCCC AGGTCAGGAG TGTTTTCTTG GTGGTTCCAG CCCCCATCAA TTGAACTGTT TCTGGGCTCA GTCAGACACA GACATTCATC TGTGTCTGAC CAAATCAGGG GCTTTCCCAC CTGTGGGGGA GGGCACAGTT AGGATGTTTT T

AGGECTIAC AAAACACAAA TITATTATCT TACCATTCIG TGAGTCAAAA TITCCAAAATA GGIGICACTA GGCTAAAATG
AAGGACIGCA TITMINCCIG CAGGCTCCAG GAGAGATCTA TGICTIACTC TITMICGGCTT CTAAAGGCTG CCCACATTCC
TCGACTAGIG GCGICCCTCC TICATCTCTA AACCCAGCAA CAACAGGITG AGTCCTCATG TCACATCTTT NITACCTITTC
TGICATCTCA TCTCGCTGAC TGCTGCTGGG AAAAATTCTC CACTTTTAAG GGCTATCATG ATTAGACTAT GCCCACTAGA
TAATACAAGA TCTCAGATCC CITAACTTCC ATCACATCTG CAAAAGTCGC TT

SEO ID NO:1034: (Length of Sequence = 320 Nucleotides)

CGCGCCGCGA CGGACGCCCT CAACCGGCAA ATCCGCGAGG AGGTGGCGAG TGCAGTGAGC AGCTCCTACA GGAATGAMTT CAGGGCATGG ACGGACATCA AGCCTGTNAA ACCAATAAAG GCCAAGCCCC AGTACAAGCC CCCAGATGAT AAGATGGTTC ATGAGACCAG CTACAGTGCT CAGTTCAAAG GAGAGGCCAG CAAGCCCAACA ACAGCTGACA ATAAGGTCAT TGATCGCAGA AGAWTACGCA GCCTCTACAG CGAACCCTTC AAGGAACCCC CAAAGGTGGA AAAACCTAGT KTTCAGAGTT TCAAACCAAA

SEO ID NO:1035: (Length of Sequence = 375 Nucleotides)

TTTTTTTTT TCAGIGGAAA ATAACITINA TIGAGACCCC ACCAACIGCA AAANCIGINC CIGGCATTAA GCTCCITCIN
CCTTIGCAAT TCGGICITIC TICAGIGGIC CCAIGAAIGC TITCINCICC TCCAIGGICT GGAAGCGGCC ATGGCCAAAC
TIGGAGGIGG TGICAATGAA CITAAGGICA ATCITCICCA GAGCCCGCCG CITCGICTGC ACCAGCAAGG ACTIGCGGAG
GGIGAGCACC CGCITCITGG TICCCACCAC ACAGCCTITC AGCATGACAA AGICATIGGT CACTICACCA TAGTGGACAA
AGCCACCCAG AGGGITGAIG CICITGIMAG ATAGGICATA GICAGIGGAG GCATT

SEO ID NO: 1036: (Length of Sequence = 304 Nucleotides)

CICIATGICT TCTTCTTTT GCTTCTCCTC AAGTAGAG I TGACTTTTTT GAAGGTTAGC TTCTTCTAAG AGTTGCATGC
TATTNCTGGC TCTTACAATA GCCTCATATC TCTNATTTNC TAATTCATTG CACTTTGCTT GTACCTCTCT GGTCTGTTTT
TCCAGATGTG TATTINCGGN TCTNAATTGG TTGGCTTCTT GGATTGTCAC ACATAATCTT ATTTCTAATT GTTTTATACT
AGACTGTAAC TGCTGTAAAC GGCTATCTGA TGCTTCCTCT CTTNCATGGG CAGACACCAC ATCC

SEO ID NO:1037: (Length of Sequence = 341 Nucleotides)

CTATGAGGAC CAGCAATTAG ATTTTATAGC AGTACTTCCC ATTAAAGTGA ATAACCAAAA TCACTTTAAG GTCAAGATCT
TAGTCAATAC ATTATGTAAA ANCATATACA ACAGACAATA CACCAGAAAC TAAATCTTTT GCAACCTTTT AAACTTATGA
TGAAAAACAT TAATGTCAGC TCTAAAATGT ATTAAGCAGT TTTTACAAAA AAAATGTATA GAATACAGGA GCCAAAACAT
TTANCAATTA CCCTAACTTG CTGACACAGA NTACTATTAA TAAATAATAC TGATCANKIN AAAGTAATCA ATTTGAAAGT
GGTGGGGGTA GAAGGACAAC A

SEO ID NO:1038: (Length of Sequence = 281 Nucleotides)

GGAGGCTGAG GTGAGAGGNT CCCTTGGGCC CAGGAAGTCA AGGTTGCAGC AAACAGTGAT TGCACCACTA CACTCCAGCC
TGGGCAACAC AGCAAGATCC TGTCTCAAAA AAAAAAAAA ATATCAGTAT TGTTTTATTA ATTGTAACAA ACACACTAAA
TAAATGTAAG ATGCCAACAC TAGGGGAAAT AGGATNTGGN GTAAATGGGA ACTCTCTGNA TCATTTTTGC AACTTTCCTG
TACATCTTAA ACTATTTTAA ATGNTTCTAC AAAAGTTAAC A

SEO ID NO:1039: (Length of Sequence = 246 Nucleotides)

CCAATGATGG CAAACATGAG GATGGCAAAG AAGAGAAGCA GCCCAATCTG CAGGAGTGGA ACCATGGCCT TCATGATGGA CTTGAGCACC ACCTGCAAAC CTGGGGCCAG AACAGGGCAG GTCAGGAAGC AACGTGGGCA GGGTAGGGCA AGGAATTTNG TOGGGGCAGG GACAGANCAG CAGGAACCTA GCAGGGACAG CAAGGTGCTA AGCAGTNAGT GCTTTCAAGG GCAAAGGTTA GAGCTG

SEO ID NO:1040: (Length of Sequence = 399 Nucleotides)

GAGGICAAGA AGAGCITAAG AAAATATAGG AGATACTACA GCATGITIGG TICATGACCG GAATGATTA GTAAGAAGGA
AAAGCCAATA ATGIAAGAAA GGCGATTGCA GGAGCAAAGA CITTAAGGAA TAAAAAAGGAC AAAATTGITT GITTCTCAGG
GAAGTAATGA CAGGGGCTGA GCAGGAGCCA GGAAACCCAG CITTTAGCTT CAGNICTGCC TGACATTTAT TGGTCATGTG
GCTCTGGGTG TATTCTCACT TCTCCTCCCT AAATAGCAAG AAGGAAAAGC CTCTTGGAGC CTCGTGTCTC TGCTTCTTTC
TGTACAATGG TTATGTTTCT GNTCCGCTTA GCTGGTTAAT TATAGAATCA CCCTNGCTGG GGTCTTTTGG GGACTGGCC

SEO ID NO:1041: (Length of Sequence = 324 Nucleotides)

CCATAAACAG TCCGTCACTG ACAAATGTTG TTACGCAGCA CATTTTATGC AGTGTGTGAC CATACACGAT ACACAGAGGA
AATTCAGGGC TTCTAGGAAA CCTTCTAAGG CCTCATCTCC CTAAGGGCAC CTGATGAGCC ATTCCTCACC CCTGCACTGC
ACCAGGACTC CAACACCACC ACCAAGGCTA ACCGCTGTGC ACTCTGGGCC CTGGGTCTGC AGTACCTGGC TCCCAAGCAC
ACCAGCATCT GAAAACTTGN CATCCTTGCC GATNITNCGG GGAGTATTGG TTGATTGCAG TGACAAATCG GCAGAAGTTC
CGGG

SEO ID NO:1042: (Length of Sequence = 212 Nucleotides)

ATCIGITICI CAGAGATGAC ACTGCCAACA ATCACAGATI TGCATACAAT ACAGITATGI ATTGGCTATT CACAATTTAC
AGIAGIGITI TICCCICTGA AAAATATAAG INCAAAAGCI AAGIAAACAA TGNGGTACTG CCATTTGGGN TITTITTACAT
GGNCTTAGCI TAAAGAACTG GICTTTAGCA AATATTCAAC AGNTCAACCT GA

SEO ID NO:1043: (Length of Sequence = 329 Nucleotides)

ACTIGEAGAA AGAAAAATTA GAGAATTCCA GATCCITAGA AIGCAGATCA GATCCAGAAT CICCIATCAA AAAAACAAGI TIATCICCIA CIICIAAACI TGGATACICA TATAGIAGAG AICIAGACCI TGCIAAGAAA AAACAIGCII CCCTGAGGCA GACGGAGCIA TICCAGATGC TGATAGAAAC ACTITAAATC AIGCAGATCA TITCATCAAA ANIAGINCAG CAGCAAGATG AAGAGGGAGC TCGGCAGCTG AGAGAGAGG CTCGTCAGCT AATAGCAGAN GCTCGATCTG GAGINAAGAT NICAGAACIT CCCAGCTAT

SEO ID NO:1044: (Length of Sequence = 285 Nucleotides)

GITGAAGCTG TITINATITC ACACCCITCT GITTIAAAAC ATAGGGACTG ACAGGGAGAC CCAGGGCTGC AATCTGGGTG
GIGCIACATT TGIAGACAAG GACAACITGC TGIATITIAA CCCAGAAACA TTAGAAAGTT TGICCITGAA CITCTGGCTC
AGATTIAGAT GCATCTITGA AGTGCTGATA TITGGCTTAT CTGAAGCTTT GGGATTATCA TTINCIAGTT ATGAAGGGAA
TGAAAGTGTT CATAACATTT TTGCAGGTGG AAGGTAAAGT TGITG

SEO ID NO:1045: (Length of Sequence = 317 Nucleotides)

TOGGITACIG TAGIATIGIA GIATAGITIG AAGICAGCIA GIGIGATGCC TCCAGCITIG INCITITIGC TCAGGATIGI
CITGGCIATA CAAGGICTIC TITGATCCCA TATGAAATIT AAAGIAGITI TINCIAATIC TGIGAAGAAT GICAATGGIA
GIITCATGGG TATAGIATIG AATCIATAAA INATITIGGG CAGIACGGIC ATTITCATGA TATTGATTCT NCCTATCCAT
GAIGATGGAA TCITTITCCA TITGITTGGG NCTTCTCTTA TITCCTTGAG CAGIGGGITT GIRCTTTTTG GACAAGA

SEC ID NO: 1046: (Length of Sequence = 316 Mucleotides)

CCAGGIGCAA TCTCGGCTCA CTGCGACCTC TGCCTCCGCG TAGTGGGACT CCAGCTGTGC ACCACCCAGT CAGCACCCACG
CCCACCCTGC CAGGGCTGTG CACGGTTCAG CGTCACTTTA CAGATGAGGA AACTNAGTCT TTGGGAAGCT GACAAGGTGC
CTGACACAGG CCAGGGCAGG GNCCACCCTC ATGGGCTGTG CTGCAGCCTC TGCCTCGTGG GTCACGGCAC CCCATCTACG
AGGNGCCCCT CAAGGATGCG CCGTCGAGTN CCCGGGGCCC TTGGCATGTN CCTGGCAGAG AAGGCAGCTC AGGGGT

SEO ID NO:1047: (Length of Sequence = 261 Nucleotides)

CTTCCTCAAA CTCGGGTTCC AGCTGGGTCT CAAACTCAGG CTCCAACTGG GTCTCAAACT CGGGCTCCAC CTTGGTCCCA
AACTCGGGCT CCACCTCGGT CCCAAACTCT GTCACCACCT CTTINIAGGT CTCANICTCC GACTCCTCCC AGCCAGCGGT
GGTTGGCGGT ATNAGGCCCC AGGGCTCTAT GGTAGTGCTC AGGGTGNGTG GCAGGGGCAG GGGCCAGCGT GGGAGGCCACA
GTGTNGGGGG CCTAGGGTGG T

SEQ ID NO:1048: (Length of Sequence = 390 Nucleotides)

GAGAACAAAG AGAATGGAGG CCACATACAA TGGAGTAACA GAAGCTTTGC CTGTAGCTCA AGAACCAAGC CGAGAATCCA
CACCTCCIGA TTCACAGITC AGIATTTCG GCCACITTAC TCAAATATT TTATAAATTA TTTTTAAATC GGCAAAATAT
TTAAATTTCA TCCATTAAAT TTAAATTTCT AGATGCCCTA GIGGCATCCA GAACACATAT TINGGGGAAA ATATTCTAAT
TTTTTTAAAC AGAAAAAGCT AGGNCAGAT GATGCATTAA AAAAGTAGAA CACAGAGCTC TTAATTTAGG AATGATCAAA
ATAGGGITGA TTCAACTATT ACCTTCTCCT AGGGATTATG GATCAACCCC TAGCAGCAGN CAAAGTCACA

SEO ID NO:1049: (Length of Sequence = 335 Nucleotides)

AAACTCACAA GIAAAATAAT GCATATTTAA GGGAAATATT ATACAGACIT TITCACACAG AAGTACATAA TANGATITIT
TAAAATCIAT TGCCATTCAT TTATTITTIGC ACAAAAACGT ATAAATATGT CACCAGCTIT NCITAACTTA AAAAACTTAA
ATAAAAGACA CCAGATGAAA ACTACCCTIT GCTGCCATTT TTTTTTAAGT TTTTTTGTAG GGGTTTTTTA TTTTTGCRGT
TTTTTTNCTT TINCTGCTTA GAATTGGGTT TCTAGGGAAG AAAAGCCCCT GCATTAAAAA CAGNCCATTT AAAAAAAAAA
TTCAAAAGTTC TGGAT

SEO ID NO:1050: (Length of Sequence = 265 Nucleotides)

AAAGGAGGG AGGGAGGAT GTGGAAAATA TGCAAGATAA ATTAAATNCT TAGTTAAAAA AAAAAAAAG TTTCACCAAC
TGINCICCAT TACTGAGAAG CCCCCACACT GCCCCACTGT GCATATTCCT AGTATTCAT CCATGTCCTG CTCGCTGTG
CTGCCCTACA AAAAANCCCT CCCGGGGGG AAAAAAAAANC AAAAAANCGG TGTAGTGTGA ACTGCTGAAG AACTTAAATG
TTCAAGAGCA TCTTTAAAGT CTAGG

SEO ID NO:1051: (Length of Sequence = 298 Nucleotides)

ATTICIAAAA TECTOTOAAA TACTAATATT ATACATTOTO COATTTATOO TOAAAAAACC CATGAGACTG GTGATGTAAT
TNOTGTGTTC ATTICACAGC TGTGGCAGTC AGTOTAAAGA COAAGTGATT TGCTCAAAGT CATGGAACAC TTAAATGGCA
GAGCTAAGGC TTAAACCCAG AATTTAAAAA TTTTTTTINAG CTTCINGTTT TINCCATTAT ACCAGTTTGG COCTTCATTT
TATTCATGGG TTAAATTAAA TTATGGTAAC AAAGGGCCCCC TGGTCACTTT GGACATTT

SEO ID NO:1052: (Length of Sequence = 359 Nucleotides)

AAGGCAAACG TGGTACATCA TGACACCATG GGAATGACTC ATGCCAGCCA TAAAAAGAAT GAGAATTCTG TCCAGAATTG GTTCCTTCCG GTGGGTTCTT GGTCTCGCTG ACTTCAAAAAA TGAAAGCCAT GAACCCTCGT GGTGAGTGTT AACAGTTCCT TCAAAGATGG TGTGTCCGGA GTTTNTTCCC TTNCAGAATG TTCCAAAATGT TATCCCAAGT TTCTTCCCTT CTUGTGGGTT TGAATGGITA ACCAACCCCT AGGCTACCAC TCTGTATTIC ATCAGGGGTA GGGGTATTAA ACCCCACATG CAAGTAAGGA ACCCTTGCCC CCAGTGTGCA AATGGGATGG GGATGCT

SEQ ID NO: 1061: (Length of Sequence = 206 Nucleotides)

AGAAAGTAAG ATTCTCAGGG CAACAGTGTA CAGCAGAGTG GTTGCTCCAC AGACAGAGGA GGGCAGAGTG GCCCAGAGTA
TCAGCGTACA GCAAAGTGGG TGTTCCCATC CACAGGGGCA GCGCTATCTC ATAGGANAGA ACAACCCCTA GGAAGGCAAG
CGTCAGNCAG NCAGCAGTGN AACAGTCAAC AGTTAGCCAG TGTCAG

SEO ID NO:1062: (Length of Sequence = 316 Nucleotides)

TINCCCICAC AGAGITITAG TIAGAATCAC TITCICIAIT TCCACAAATC CITCITITCT TICCIITIAT TITCIAAAGI GAATGICCAA GCAAAAAGGA AGCAAAAATG GICAAAGATC TCTCTTACAA TATAGIAATA AATITAINCA AACAACTIGG AATICACCCT GIGCATIGAA AATINCAACTC CACACTGCAA ATTATGGCAT TITTICCCNC TCAAAGGAAT TAGTGAACTC CATTGGATGC ATTCATACIN CIGITTAGGN AATAAGGGAA ACCGCTTTGT AAAAGINCAA CATGGCCTAG GAGTTA

SEO ID NO:1063: (Length of Sequence = 314 Nucleotides)

ATGATCTGGT TTATGCTTCA GAAGAAGCAT AGTAGCTTCT ACAGAAAATA AATGATAGAA GGCAAAAGAG AAACATGGCG
AGTATTCCAC TCCAGTGTCT AGTCAAGAGA TTACAAGGGC CCTGGCATGA GGACAACAGT AGAAATNGTT AAAAGTGTAC
TGGATTGCAA AATATTACTT TTGGGCCAGG GCGCCGGNGG ...ACACGCCT ATTAATACCC AGCACTTTNT GGAGGTGCAG
GGAGTTNCGA GTACCAGTCC TGGGCCAACA CGCNTGGAAA TCCTGTTGAA AAATATAAAA ATTAGCCGGG CCGT

SEO ID NO:1064: (Length ci Sequence = 322 Nucleotides)

GAAAGCATTI GAACTAAGIN TGTAAAAATG GCAGATAATA ATTAACACTI GGTAGCAAGA AACGCTITCI GAAATACTGG
GAACACTGAC TIGITTCACT GTAACTTATC ACCTAGTGCT GTATCTGCCA TAGTGCTCAC AATTGCAACT TTATATCCAA
CATGGGTGTT CCATTTCTAT TIGGATAAAA TTTACTGGAA ATATACTAGC AANGAAAAAC TGGTCTTAAA ATGGCAAAAG
GCTCTGGCAC TAAATTCACT GCTACTTAAC TTAGTTTACT AATTAACTTC CTTAATTATA GTTTTCCAAA TCCGCATGCA
CG

SEO ID NO:1065: (Length of Sequence = 297 Nucleotides)

CCCTENCAAC TCCTTGCATG GACTGATGCT GGAAACTGGG TCAGGGAGCT CCAGGAGGAA CCAGACAGGN TCCTGTTAGC AGGCTCACCA CAAGTTCTAA AGGGCACCAG CCTTGAGAAG GGCAGTTGGG ATGTGGCCAA ATGTGAAGCC AGGTTTNCTG GGATCCTGAC TGTCCCAAGCT TACAAGTTCC TGGCCACTCT GTGAACCTTG GGCAAGTTAA CTTCCAACCT CTTTACAAGT TCCCTAATCT ATNAGGAAAC ANTTAGTNAC ATGACCTTCA TGGGAATTTA TTTATGA

SEQ ID NO:1066: (Length of Sequence = 267 Nucleotides)

ACAATGGEAC TGTCAGAGCA GCCAGCTCCT CCCTGACTGC TCCACAGGAA GAGCCATCAA CAAAGCCAAT CCCTGGAGAT
AGGCTCTGAA ACCAGGATAG AGACTCCTTC AATGGCTGCT GNTGGTTCCA CCATGTATCA TCCAGAGNAA TCACCCTGNG
TGGGCATAGG TGGGCCTGGG AATCTAGGGC ACAGCAATTC CACACATCTT CACCTAGAAA CCCTCCTTCT GGGTGGGCCT
GCATGGTTTC ATGCCTGTAA ATCCCAG

SEQ_ID_NO:1067: (Length of Sequence = 220 Nucleotides)

AAAATGCAAT TGGTTTGTTA CTGAGTACTA TTCGTGGGAA GACAGCATCC TGNACTCCCT CTCTACAGAA TATTGGGAGT AAAAATGAAT GTCATCCCCG GTGGGAAATA TTATTGGGGG TTGGAAGCAC AGAGCACAGG AAAAATTAAG TNCAGGAAAC

281

AGACACTAAG AGTGCACTGG GCAGGTCTGA CTGCAGGTGA TGCAACTTGC CAGCCGTGGT

SEQ ID NO:1068: (Length of Sequence = 412 Nucleotides)
TGGCCAGCAT CTGGGAACTT TGGGTTGTTG GACCAACTTC TTCCAACACG TGCGCACTGA TGGCCGGGC CCCAGCCAGG
CCTGNCTGGA AGGGTCTTCC CCGNCCCGAG GGACTGTAGG GGGTCTCTAG GAAGCATCAC ATCAAGGTCC TCAGGTTAGA
TNCAGGNCAG CCCATTGACC CATTTNAGGG GACAGCTGGA GGGAAGCCCG GAGTCCCTTG TTTCTTCAGC TGAG
234

TOSCICATGA AGATAATITA ATGCTAGACT GATTICTGCA GAGTAAAATC TGGCATGINC TICAGGAAGT TITCTITGTC GCTGCATATG AAACATTAGG TCTCCTCCAT TTACATACTC TATAACAAAG AACAATCTGC TTTCTGTCTG AAAGCAAGAA TGCAGCCTAA CAAGGAAAGG ATGATTGGAT GCCTGCTCAA ACACATGCTT CTCTGTCTGT ACCCAATCAA TATCCTCATC ATCATTAACA AGCTCTTTTT TCACAACTTT CATTGCATAA ATACGATCTG TTTTTTTTTAA TCGAACCAAC AGTACTTTGG CATAACTTCC TCTTCCTATT ACCCGGAGCA AATCAAAATC CTGAAGACCT AGACTGGATG AAGCTTTGCA CTTTCCCTGG NGTCATTGCC TC

SEO ID NO:1070: (Length of Sequence = 358 Nucleotides)

SEO ID NO:1071: (Length of Sequence = 411 Nucleotides)

CTATTTATGA ATTCTOGCAT TGGTTTOGAA AACTCAACAC AGITAAATGA ACAGGAATTG AAGGTGCATG ATGGATGCGT
CCCTCATAGC ATTTAAATCT CTTCCACTTG ATTAAAAATT CCTAGTTCCT CTTCACTGAA TTGTTTAGAG TTTTTNAGCA
GCCTCTGCCC TGATTAAAAC AAATTAGCAT CAAAGATCCC CTGTTGAATG AGAAATCATT AATTGAGAAA CATGCAATGC
TCCTTAATTA CTTTTAGAAC AGTGAGAGAA CAAATAATCT CAGGTTCCAG AGGGCCCTGC CTGCTCTGCA CCGTGAACTC
ATTTCGTGTA GCTGCTGGAA TAAAACTCAA GTAGGCAAAC ACTATTTGGG GAATATCAAT GCAAGCTTTC AGTAAACACA
CTGTAGGATT G

SEO ID NO:1072: (Length of Sequence = 342 Nucleotides)

TCCCATTITI ATAATIATIG GAACAIGAAA CIGIATITCI ATGAACICAA TGAITITITIT CCATAAAATI ATATGCTAAG
AGAGICACCA CAAAACTAIG AATICICICC CGAATIATIT TIGCIICITIT GGAGCACCAT AGICITIGIT CAAATCACAA
CAIGAAACIG TIGCIGCAAT GCIAAAGAIG TGAATCCACC ACTATCAATA CGGICAGGGI AAAACCIGGA GCCACATGIT
ATICAAGITA TITTIGITAT CTAATGAITG ACATGAAAAT AAAATAGIAA GCCAATATIA AATITGIAGG CATAGITGCC
CCACCINAAA AGIGITITACA AA

SEO ID NO:1073: (Length of Sequence = 217 Nucleotides)

GITTCIGIC CIGGCTAGGA TAATGCAAGC NCTITICAGA TGANICAGAA TOGAAGAAAA TACGCIGGIA AAACAGGACC

TGATTCACCA GENACTAAAC AATTACACIC CCATTICCAT TGCTTCAAT ATTITCACAC GNIACACAA CCTTTAAGAT

GGAAAGGGAA AGCGATTITT INITCAACAA GTGGGCCACC AGATGAACCA AATTAGA

SEO ID NO:1074: (Length of Sequence = 379 Nucleotides)

GGTTAAAATT TCATCGGAAT GTATAAGCTT ATTTATTAGT GTATTTAATG GTTCATCAAT TGATAAAACA GGTGTAGCAA
ATACATGCCT TCCTTTTGGG GGATGGGCCT GGTTAATCTC CAAATTGGCC GTTTGGAACA ACTCATCATT ACTGTACAAA
GAAGGTACCA CTTGGTGGGA ACTTTCACTT TTTAACAAAA CTGGTTCATA TTTCTCACTT GCATAGGAAA TGGTCAAACC
TTGAAGTGAA GCAGAGTGCA TATGAGAAGT AGGCGACACA TCAAAAAACTG GTACAGATGT AGAGTGCAGC ATGTTTTCAC
TTGAAGCAGA ATTTGATACA ATGAGGATGC AACCATTGTA GANCTAAATT TATCAACTT

SEQ ID NO:1075: (Length of Sequence = 345 Nucleotides)

ATTAAGITGA CAGICCAATC AGAAATATIT AAACAAAGIT TCACIACITA AACACCATCI AAATATACIT TITGITATAT TCCCAGCAGA AATTGATGGC AAGGAATCAT ATATCCCATC AAAACCGTAT TTTTCCCCCT AAAAGGCAGI TTAGATGINC TCATTCTAGG NITTCCATCI CICTCCTCCA CCATTCCAAT TCCCAGAGIA CCTCTACAAA TATCCCTGCT TACCAGTAGA NCTATTTGCT TTAACAATCI TICTGTGGGT AAGGAGATGC ATATGCCAAT GTGAAAACTA TGGAGGGGGA CTCCTGCCTT CAAAGGCTGA CTAGAAACCA TTGGA

SEO ID NO:1076: (Length of Sequence = 286 Nucleotides)

TTTTTTTGA GATGGAGTCT CGCTCTGING CCCAGTGIGG AGTGCAGTGG CATGATCTCG GCTCACTGCA AGCNCCGCCT
CCTGGGTTCA TGCCATINTC CTGCCTCACC CTCCCGAGTA GCTTGGACTA CAGGCGCCTG CNACCACGCC CAGCTAATTT
NTINIGIGIG TGTTTTTGGC AGAGACAGGG TTTCACCATG TTGGCCAGAA TGGTCTCTAT CTCCTGACCT CGTGATCCAC
CCGCCTTGGC CTCCCAAGGT GGTGGGATTA CAGGCGTAAA TMACCG

SEQ ID NO:1077: (Length of Sequence = 366 Nucleotides)

TCACATAGGT CACATTITAC CCATGAAACC TITCTAAATT ACCITITGCA TITNITGCCT ATCCITCTAC ATCATCATAC
TTCGTCAATT AAAGTCACTT TITTGGGTAA CATTTCAGAA ATTGGGATTC CICTTACAAT TGCTATCAGA CAGAAGCCAA
TTATGATGTT GTCATTGCTT ACACATGGGA AAATAACAAA ACTGCCAGCA TGACATTTGC ATATGACAGT CAACAGCCTG
AAAGAAATTC CCAGAAATGA TACTGGAGCA TTCATTTCAC CCTCTAGGAN CCAAATGGAC TNGGAAGGAA GTAGAAGATG
GGGAATCCCT AAGCAGCAGT CAAAGTAGGC TGGCTTTTCA TAATTT

SEO ID NO:1078: (Length of Sequence = 380 Nucleotides)

GITTAAGIGC GAAGATITTA TIAGGOGGIA CAATICCAAG GIGGIAAGGG TGAAAGGAAA GGCGAAGGCA GGCAAATACA
TIATIGAGCI GAAAACAACI TIACATICAA GGACAGCTIC CAGACAAGCC ATGIAGAACC AGCATGCCIT GGGACTGINI
GGATGGCAGG GAGACGAGTI TCIATGCTGA CCACTICATG CTITCISCCC CCTITGGGGA AAGIATGCCI CACGGACCTC
TAACTCTCCC ACTICICIGG GGGCAGCACC TGACCCCTCC CGGCAACTNC TAGGCAAGAG CATTCTGTTC CTTCAAATTT
YTCACCTGAG TCIGAGICAG AGCATYCCAT CATCAGAGCC TCTGTCAAGG AGGCAGTGCT

SEO ID NO:1079: (Length of Sequence = 439 Nucleotides)

CTTAAGTTAC TGAAATTGAA ACACCCITTG TCCTTCTCGG CGGGGGCTTC CTGGTCTGIN CTTTACTTGG CTTTTTTCCT
TCCCGTCTTA GCCTCACCCC CTTGTCAACC AGATTGAGTT GCTATAGCTT GATGCAGGGA CCCAGTGAAG TTTCTCCGTT
AAAGATTGGG AGTCGTCGAA ATGTTTAGAT TCTTTTAGGA AAGGAATTAT TTTCCCCCCT TTTACAGGGT AGTAACTTCT
CCACAGAAGT GCCAATATGG CAAAATTACA CAAGAAAACA GTATTGCAAT GACACCATTA CATAAGGAAC ATTGAACTGT
TAGAAGGAGTG CTCTTCCAAA CAAAACAAAA ATGTCTCTAG CTTTCTCTAG ACGTAATTACC CTTTCTCTAT —
TNAAATCAGG GTAACCCCTT TCTGTATTTG AGTGCAGTG

SEO ID NO:1080: (Length of Sequence = 419 Nucleotides)

CTGAACTCCC TGAAAATAGG AAGTCTCAAT TAAAAAATCA ATTTGTCATA GTCCACATAA AGATAATCAA TACATTTTGC
TCTCAGTCCT TGGGATGGTT TTTGTAAATA ATATTATCTT GACAAAAACA AACAGGAAGA TCCCACCCCC AACACATACC
ACATTCCAAT GTTACCTGGN ATTAAAATAT ATACCAACAT GCATCTTTAG GTTACTCTGG TCCATGGTTT CCTCCAGTGG
CAATGGAATT TACAAAAATG TAAGACGTAA TAGATATATA ATTATCTTTT TNCCTAAATG AAACTAGCCT TAAAAACTGG
TACATAATGG TTCCTGGGTT CANTGATCAA AATTATGGAN GTACACTTAA CCTATCTTCC ATTGAGTGGC TTTAAAATGGG
ACCTTAAACT GTGGACTCC

SEO ID NO:1081: (Length of Sequence = 411 Nucleotides)

CAGCGITTAA ACCAAAGGCG CACTAAACCT CGTAAGCGCA TGANCAGATT TAAAGAGAAA GAAAACTCTG AGTGTGCCTT
TAGGGTCTTA CTTCCTAGTG ACCCTGTGCA GGAGGGGGCGG GATGAGTTTC CAGAGCATAG AACTCCTTCA GCAAGCATAC
TTGAGGAACC ACTGACAGAG CAAAATCATG CTGACTGCTT AGATTCAGCT GGGCCACGGT TAAACGTTTG TNATAAATCC
AGTGCCAGCA TTGGTGACAT GGAAAAGGAG CCAGGAATTC CCAGTTTGAC ACCACAGGCT GAGCTCCCTG AACCACGTGT
GCGGTCAGAG AAGAAACGCC TTAGGNAGCC AAGCAAAGTG GCTTTTGGAA TATACAGAAG AATATGATCA GATATTTGCT
CCCTAAGGAA A

SEO ID NO:1082: (Length of Sequence = 350 Nucleotides)

CTGTGAGGGC ACAAGTGTAG GTATCTTINC AAGTTCTCTA GGTGATTCTA GAATGCAGCA GGGTTGAGAT GCTCTGCCTT
AGGGGTAGAG AGGTGGGAAC ACTGACAGGT TCTGCAAAAC ATCTCTGAAC AGCTGCTGGT GTCTTTTTCT GTACTTCAAG
TTTCACGGCA CATCTGATAG CTGINCCGAA AGGGAAGAGA GAATTACGTG GGCTAGGCTG GTTTGAAGGT TTGCNTAAGN
TTTGGCTTGA GCGACTTTAA CACGTTTTATT TCAAAGTAAT TTGTGTTTGT AGCCCCACTA AAGTAATTTT GGGCCAGNAA
AGGTTCAAAA TACGGTTTTC CCTACTTAAG

SEO ID NO:1083: (Length of Sequence = 430 Nucleotides)

SEO ID NO:1084: (Length of Sequence = 369 Nucleotides)

AATGGAAGAA GTGAAAAAGA ACAACACAAA GAAAATAAAG AAGTAACCTC TTTCACCCAC TGAAATAATC TCTGGAAAAG
ATATTAGCAA TCATGCAGCT TATAAATATC TAAAGGGCTA GAATTGAGGA ATTTATAAGA MTAAMTTTT TTTTCAACAC
ATAAAATACA ACATGGGAAA TAAGATGTTT TTTACTAACA GGCAAACACT TGAGGAGTCC TCTTCAAAGA CTACAGTGGA
TGAAAGACCA GTTATCCAAA GGAAACGGTT AGTAGAAATA TAAAGTTAGT CCCACACAAA ATTAAAATGG TGCTCAATGC
AGATTATCTA TCATTANACC ATTTTTAAAG GCAATTTNTT ATTTAAAAT

SEC ID NO: 1085: (Length of Sequence = 413 Mucleotides)

ATACCTTINA GCIGGCATAA TITAACGTIC TAATTATCCC TIAATCATAA GCIGIACGAT TCIATAATTA AAAAGITAAT GCCITCITAA TGICIAINCI AGIAGAAGAA TGATGAGAAA ATAATAGIAT AGATTAGIIT TGGICICIAC TCATTITGCC TTCTGATTAT ATTACAACTC CAGCTGGTGA CAAGATGGCT GTGTAAATCT TGAAATCACT GAGCATTCAT TITAGCTTCT CATTGAAAGG TAGATATTCA GTATGAATTG TAAACTGGCA TTAAGGGAGA AAGTAGGAAT AATCAAACTT GATCTGAGAA TTACTTGCTG GTGCATTTCC TCAATGCATA GTAATATCCT TATGANGATG CAGATGCAAA AGTGGGTTTTT GGAGGTGGAT AAGGAGGGCA GCT

SEO ID NO:1086: (Length of Sequence = 277 Nucleotides)

SEO ID NO:1087: (Length of Sequence = 360 Nucleotides)

TTTTTTTTT TTTTTTGAG ACATTGTCTC ACTGCGTCGC CCAGGCTGGA GTGCAGTGGT GCAATCTTGG CTCACTGCAA
CCTCTAAATC CCAGGTTCAA GCGATCCTCT CACCTCAGCC TCCGGAGGGC NTGGGATTAC AGGTGTGAGC CACCGCGCCC
GGCAGCATTA TTTTTTAAAG ATCTGTGATA GTGCATGTTG TGCTAGTTCT TTAATACAGA CTATATTGTA TTCCATGTCA
GTTTTTAAAG TTTATTTCCC TATTGATGGC ATTTAATTCC AACTTTTAGA TAAAAGGATG TACTGGACAT TTTTATAATT
TTTTTGGGGG ACCATGTAAG AGTTTTTCTA GGGGGAATTC

SEO ID NO:1088: (Length of Sequence = 209 Nucleotides)

CTGGGACCAG CTGGAACAGA AGTGGTAAAG GATAACTAGC TACCTGCACC GCCAGAGATC AGGNTCAGGG TGAAGCTGGT
TTCCCAGCAG GCGAAGTGAA GGAAAGTGGT TNGAAAGGAA GAGGAGGAGC AGGAGATGGT AGGTCCCTCG CCTNTCTCCC
MINCTACCCT GGAA:NATAA GTGTCAGGTT CATACTTAAC CACCCCCTT

SEO ID NO:1089: (Length of Sequence = 409 Nucleotides)

TITTIGCTCAC AGCTACATCT TCAGAGGTGA GAACCATGCA TGACACAGAG AAGATGCTCA CTGATGGATT TAATGAGTCA AACATTGAAG AATCAATGAG TGCCGGAAAT AAACAGGATA GGTGGCAGCA TAGCATGCCC TTAAGANCAT GGCTGTGGAT TCAAATCCCA GACCAATCAC TGANITTCAA GCCACITTIGC CTCTCTGAGC CTCTGTTTTC TCATCTGTCA AGTGGCAATA ACAATAAATG GTACGTGCCT CATAGGGGCA CCTTGAGGAT TAAAAGAGAG GGTTTCAATA AATCAAGTAC TGATTTCAAA ACCTGGCACA TAGTAGGCAC TCAGCACATG GNCCTTATAT ACTINTGGGC CAGCAGCGGC TGGGGCTCAT CCCTCCCTGG CTGGGTCCA

SEO ID NO:1090: (Length of Sequence = 337 Nucleotides)

GAACCINTCC CCATTGGAGA GGATGAGGAT GATGATCTGG ACCAGGAGAC ATTCAGCATA TGTAAGGAGA GGATGAGGCC CGTNAAAAAG GCACTGAAAC AGCTGACAAA ACCTGACAAG GGGCTCAACG TGCAAGANCA GCTGGAACAC ACCCGGAACT GCCTGCTGAA AATCGGAGAC CGGATAGCCG AGTGCCTTAA AGCCTACTCA GATCAGGAGC ACATCAAACT CTGGAGGAGG AACCTATGGA TTTTTGTTTC CAAGTTTACA GAATTTAATG CTCGAAAACT GCATAAGTTA TNCAAGATGG CTCATAAGNA AAGGTCTCAA GAAGAAG

SEO ID NO:1091: (Length of Sequence = 411 Nucleotides)

CCACIACCAC AGGAAATCTC TATACCCITC TIGGCTTTTC CITTTAATGT AATTITCITA AAAGCTTCAA GATAATTTTT

AATCAGGCAT GCTGAAATCT ATCTAACCTA TTAGTCACTA ATTATATTCT TCAAGCCTAT ATATTAATGT TTCTNCTGTT

GTAAATTCAT GATCATAAAG TTTTGGACCT GGCCATCAAT ACTAAAGCAC TGATATTTAG TTTTTAGGTGA TACTTGGGCA

285

TAAATACAAA CACGGGATAT ATTINGTCAT AGAAAAAAAT GTGTTACTGC ATTATTTTGC ACTICTGAAG GACTGCAAAC
ATTITTCAAG CACAATAAGC AAATTCTTCT TTCAAAAAAGG NATACTTING CACATATGIN AGGITTGGAA AATGACTAGG
NCCCTAGGGA G

SEO ID NO:1092: (Length of Sequence = 349 Nucleotides)

AAAGAAAATG CCTTGGGAAG ACAGATGCAT TITTINCCCAC TGGTGTTGCA ATTGCTCAAA TATTTINAGG ATGAATATCC
TCACCTTGGA GGCAAGTTIT TAAGAGTGAA
TTTGAATTAC TGGAGCAGTG AACAATTATT TAGAGTCTGG TATAAGTGAA
GAAAAGAATC ATGACCNGTA AGCTGTCTTG NAGGTACCAG CAAACTGNCT CTAAAATTTA TATGGAAAGG CAAAGGGGTT
AGAATAGCCA ACATAATACT GNAGAAGTTG GAAGACTCAC ACTATCCAAT TTCAAGGTTT ACTGTAAAGC TACAGTAACC
AAAGGCAATGT GGCACTGGTG AAAAAGTAA

SEO ID NO:1093: (Length of Sequence = 400 Nucleotides)

GGACCTIGIT TIACATICIG GATITICCIT TITACTITIC TAATGATGIA ATTIAACINC TICCIGIATI TNCCATATIT
CCTATAAAAT GGIAGTIAGA TCIAAAAGCT TGATTIACIT ATTICAGATT TCTAGTCAAG GGIACTCAAT AGATTGIATI
TCCTTTIGCC TCACACGGAG GIGCATAATG TCIGCCIGGC CIGTAGTGAT GCTAAGGTIG ATCATTCIGT TCAGGTGGCA
TCAGTCIGIG ATAACTICCT GIAAGAATCG TICATTAACC TITCATCIAA TGGNICCATT CATTCATGAT CITTAACTGA
ATCCCTGTTA TITCATTAGG GAATAGCAAA ATAATGATTT TCTAATTCTG TNATTCCTTT CACATTTATT AACTGTAATT

SEO ID NO:1094: (Length of Sequence = 414 Nucleotides)

GICAGINITC CATAACIGIT TCCTGCTGAC AAAGGGGCAG TGGTGATGGT TCINIGGGTC TTGGCCTCTT GCTAGCTGTC
ACAGCAGGAG GGTGGCTTIN TGGATTGGTG AAAGTGGTAT CCAGCCAGGT CCAAGAGAGA CAGGGGCAGG GTTTINCCAA
TGCCAAATAT ACITCAGCAG TAGAAGCCAC AAGATTACAT TATTAAATTG TCCCAAGAGT CCCCCAGTGC AAACCCCCAGC
TGAACGCCAT TTAGTTATAT NCTGGTGCGT TTTCCTTCTG CAGGAACTCA AACCAAGGTT TCTTATGTGT GCTTGAGTTG
GGGGCCCAGAG TGACAACTGG TAGAAAACTA TGTTATTCCC CAGCTANGAG AACAGAGGGG AGGGGTACAT GATAGTAGGG
AGTCAAGTTT ACAA

SEO ID NO:1095: (Length of Sequence = 387 Nucleotides)

GATCIGGCAA CCAATTATGT AAATAGICAT ATGAATCCIT CAGAATGGAT AACACAGCIT TNCTGACTGG TGIGAAATAG
TITTCAGGIG CICATTCITT ACTICATTAG CITATCITAT ATCATTAGCT TATCCICCAT TCAGGIATAA CAGATCITIT
TITTCIGATA AATATGGCAG TITAGGGAAA TAAACTATGG CATAATATGC TAGGCCATTC TTCTAGGCCA CGCTTCITTG
ATTGTAACCT TAAACCCITT ATCAGAACCT AAACAACTIT TCAAAAGATC TATACATATT TNNATCCAAT GITTAAGGCT
ATGAGTAATT CATTATGGTC ACTCTTCATT TITNICACCT GATAATGATC TCGNCAAAAA TGTTGAG

SEO ID NO:1096: (Length of Sequence = 416 Nucleotides)

SEO ID NO:1097: (Length of Sequence = 406 Nucleotides)

CTGACCTCGT GATCCGCCCA CCTCGGCCTC CCGAAATGCT GGGATTACAG GCGTGAACCA CTGCGCCCGG CATGATTGGC
ATTTTGGGCT AAATAGTTTC TGTCCACAGG ACCGTCTTGT GCAGTGCAGG TCTTTTAGCA TCCTGGCCAC TCATAGTGCC
CGTGGTTCTC AGTAGAAGCT GTAGAGGATG TTGGGAAATT GGGGTGGGTT GGTCACAGTG CCTGGCATCT GTCTCAGGGT
AAGGGCTTNG GAGGCTCAAG TGCAGAGTCG TATCTGGATG CCAGCAACAC CCTGTTGAGA AACTTTCTAC TATGGTATGC
TCATCATTCT CTGAAGATGT CAGGGCCTGT TTGTTTGTTT GCCTGTTTCT CTCACTTTTG CCTTATAATC AGTTCTTCCT
TGTTGG

SEO ID NO:1098: (Length of Sequence = 326 Nucleotides)

AGCCCCCCCC CCTCCGCCTC CCAAAGTCT GGGATTACAG GCATGAGCCA CCGCGCCCGG CCATGTAACA ACTTITATAA AACACTCCC CAATCTGAAA CATGACCAT TITTCGTTTIN ACACCCAGAT TGCTCCAGAC TIGGTCAGTT GGTGTCCTC CAAGCTGGTG CTGGTCCCT TCCGCATTAGT TTTTGAGCAC CTGGACCAGT AAGGTGTTCA GTCTCACTTT GCACTT

SEO ID NO:1099: (Length of Sequence = 342 Nucleotides)

GAAAACGAAC AAGITICAGC AGICIAGCCI TIGGATGACC TATITIGAAAA CCACTGAAAG TCGIGGAGGA ATGGGCAAGA
ACCACCICAT GATICINCAG GCCATIGCTA ACGAACAGCI CATIGCTACA ACCAGICCAG AGGITITATI CCCTCTACTC
CGAGCAATGA AATAGACCIG AGITATGCIT CCTTTCATTI AATTICTGCA GATAAATAGI TICCTGAGCA ATGGATGCTA
TGCCTGGATA CCAGICTCCA CTTTGCACGC CGGAACTGCC TTGGGNCCAC AGITACAGAA AAAATGTAAA CTCAGAGTGA
TCCTTGTGTA TATTGCTATA GA

SEO ID NO:1100: (Length of Sequence = 301 Nucleotides)

ATCECTTEAG CCCAGGAGIT CGAGACCAGC CTGGGCAATG TGACAAAACC CAATCTCTAC AAAAAATACA AAAGANITAG
TCAGGIATGG TGGCGCATGA CTGCAGTCTC AGCTACTTGG TAGGCTGAGG TAAAAGGNTC ACCTGAGCCC GGGAAGTAGA
GGCACAGTGA GCCATCATTG TGTGCCACTG GACTCCAGCA TAGGGAAGGG GACTGAGACC GTCTCAAAAA AATTAAATAG
AAAGTCTTCT TTTTTTAAAA TNCTGCAATT CATGAGAAAA CTGCACTCAC ACATAGTGTG T

SEO ID NO:1101: (Length of Sequence = 300 Nucleotides)

TTAAGTCAAA GGCTAGAAAT GATTAAACTT AGTGAAGAAG ACATGTCAAA AGCCGAGAGA GGCCAAAAGC TAGGCCTCTT
ATGCCTAACA GTCAGAAATG CAAAAGNAAA ATTATTGAAG GAAATTAAAA GTGAAACAAC CTTATTGCTG ATATGCAGAC
AGTTTTAATA TTCTGGATGG AAGATCAAAC CAGCCACATT TCCTTAAGTC AAAGCCTAAT CCAGAACAAA ATCCTAACTC
TCTTCAATTC TTACGANGGC TGAGAGAGGT AAGGAAGTTG CAGAAAAAGTT TTGAACTAGC

SEO ID NO:1102: (Length of Sequence = 174 Nucleotides)

GAGATOGAGA COATOCTGGO TAACAOGGTG AAACCCCCTC TCTACTAAAA ATCCAAAAAA ATTAGOCGGG CGTTGCGGCT GGCGCTTGTN GTCCCAGNTA CTCCGGAGGC TGAGGCAGGA GAATAGOGTG AACCCTGNEN GGCGGGNTTG CAGTGAGCCC GAGATOGGGC CACT

SFO ID NO-1103: (Length of Sequence = 360 Nucleotides)

ACAAGGTCTT GCTATGTTGC CCAAGCTTGT CTCAAACTCC TGGTCTCAAG CAATCCTTCT GCCCTGGCCC TCCCAAAGTT CTGGGTATTA CAGGTGTGAG CCAGCACTCC TGGCCCATCA CAGTCTTAAA ACCAAAAGTT CTGTGTCCGA GGAAAACCAG

288

CGCTCGINIG TCCCACACAA ATGITTAAGA AGTCACTGCA ATGTACTCCC CGGCTCTGAT GAAAAGAAGC CCCTGGCACA
AAAGATTCCA GIGCCCCTGA AGAGGCTCCC TTCCTCCTGT GGGCTCTCCT AGAAAACCAG CGGGACGGCC TCCCTGCTGA
TACCGTCTAT AACCTTAGGG GGCCCTCGGG CAGGCAAACT CATCTCGGTG ATGCTGTAG ATGCTAACAC TGGCCAATTC
AATGNCACAN CTACTGGTTA CCCCTTTTGA GGGGCATTTC TCCAGACAGA AGGCCCCTTG AAGCCTAGGT AGGGCAGGNT
CAGAGATACA CCCGTNTTTG TCTCGAAGGC TT

SEO ID NO:1110: (Length of Sequence = 218 Nucleotides)

GITTINITCA TITATINNCT CCCCATAAAA CAGTATGTAC AAGGGTTIGA TICAGGGGAG AGAAAGGATA TATGAAGACA CATTCTTCCC TCTTCTATTC TCTTACCTGG TTAGAAATAA ATAGGCATAT AGTCCNGTIT ATTATGGGCA GGAAGGTAGG TAAAGATCAC CTAAGINCTT ATGGCGTGTT GGCTTTGGCA CATGGAGAAT GAGTTTTT

SEO ID NO:1111: (Length of Sequence = 211 Nucleotides)

TTIGCTTTAT GAAGAACCTG GCCTAGGTAG GGTTACAAAT GGGTTTTACT GAACTTAAAC AGCTAATTGC TACATCTCTG
AAAATAATCA GAATAGAAAA ATAGATGGAA AAATTTCAAA CCCACTGTAA GAGACTAACA TAAATCCAAT TCCAAAAGCT
GTTAATCATA CCATCTAAAA AGAAAACTGT CGACTAATCA TGTGTTTACA A

SEO ID NO:1112: (Length of Sequence = 360 Nucleotides)

CCCTATAATA GTCCCGTGAA TAGGGCTAGC AGTGGGATTT TTGTGTTATA GGCGAGGAAA TAAACACTCC TTTTGCTGAG ACTAAAGAGC CAGGTTGGGG TCTCTGGACA CATAGTGCAA TCAAGGGAGG CTTAAGACAG CAGAGGCCCT CAGAGAAGAC GTTCATTCTC CCAGCTACTT GCTAAGCACG TNCCGTGTGA TCTGGGCAGT CCTGGGCACA CCAGTGGTGA AAATACATGG TCCTGCCTGC CTGCCTGGAG CTTCTATTTT CCTNATGGAG GAATGCTGCT CCATTTTGTT ATTGGAGGAA CTTTTTGCAA GCAAAGCCTN TTTGGGAGAA AATGGCGGGC TAGAAACCTG

SEO ID NO:1112: (Length of Sequence = 448 Nucleotides)

GOGGETACTE CETTAGTGAT TAGAGITTIT NCCCTGCCGG AGGIGGGATA CACGGTAGCA TCATGGTCGA GGAGGTACAG
AAACATTCTG TACACACCCT TGINITCAGG TCGITGAAGA GGACCCATGA CATGITTGTA GCTGATAATG GAAAACCTGT
GCCTTTAGAT GAAGAGAGTC ACAAACGAAA AATGGCAATC AAGCTTCGTA ATGAGTATGG TCCTGINITG CATATGCCTA
CTTCAAAAGA AAATCTTAAA GAGAAGGGTC CTCAGANTGC AACGGGATTC ATATGTTCAT AAACAGTACC CTGCCAATCA
AGGACAAGAA GTTGAATACT TTGTGGCAGG TACACATCCA TACCCACCAG GACCTGGGGT TNNTTTTGAC AGCAGATACT
AAGTTCCNGA GGATGCCCAG TGATCAGNTG CACAGTCCTA GCGGTGGC

SEQ ID NO:1114: (Length of Sequence = 268 Nucleotides)

GGCCGCCAGG TGGIGCCATG NICTINIGIN CTGIGCGICG GGCGATGIGG TCATCAGCCT GAGACCCAGA TAGGCTGAAC CCCGACTGAT GTAGGITGCG CACAGGAGGG ACCGAGATCT TGCCTGGGCA GGACGCGCG GCCGGAGCGC CACTCCCTGG CTTGGCAGGC ACCATCACCT CGIGGACGGG CCCGTNATAC AGCCCACGGG GCACACCGTG GNITCINCGN CAGCCTGTTG CGAGCTTTGA TCCTCTTGTA GACAAAGT

SEO ID NO:1115: (Length of Sequence = 342 Nucleotides)

ATCAGIGCCI TCITCAGCIC TATCIGGGAC ACCATCITGA CCAAACACCA AGAAGGCATC TACAACACCA TCIGCCIGGN
ASTICITCCIG GGCCIGCCAC TCITGGICAT CATCACACIC CICITCATCI GITGCCATIG CIGCIGGAGACAC CCACCAGGCA
AGAAGAGGGCCA GCAGCCAGAG AAGAAAAAGA AGAAGAAGAA GAAGAAGGAT GAAGAAGAC TCIGGITCIC TGCTCAACCC

289

AAGCITTCTC CAGATGGAGA AGAGACCATC ACTGCCTGTT TAGTTAGGCA GGAANGCAGA GGTGTTTCCT TTCTGGGGCT AAAGNCTCCT TCTGACCACA CA

SEO ID NO:1116: (Length of Sequence = 416 Nucleotides)

CACCITIGGE AGGIAGGGAT CATAGITCCA CITCATIGAT GAGGAAAACT GIAGIGCAGA GATGGCATAC ACTGICCAAG
AACATGGIGG TGGATGGAAC CCAAACCCCA ACTITIGCTC CCATGINCTC TGTCCACTGG CTATGGCTCT TGCCCCTGTG
TACAGATACA GGCTCTGGAC AAGITCACCA AATCCCTTAG GCTTCAGCCC CCTCATCTGC AGAATAGTGG CITGGATTCC
ACCATCTTCA AGGICCCTGC CAGCTTINAT TTATITAAAAT TTGGATTTAT TAAGCAGGAA AAAAAGIAAT GGGAGITTGT
GGGIACCAAT GGATTAAAGG GGGINAAATC TGGAGCTCTG TGAGTAAAAT TAGGGTCCCC AAATGG

386

AAGGACCGGG ATTCTGATGA AGCCGTGTTT CTCACTGCCT GAAGTTCCC TTTGGAGTTC CAAAGTAAAG GACACATAAG CAACACTTCC AAAAACAAGG GAACAAGGTG GTTTATTGTA AAAACAGGAA ATGGTGCATG TCATTGAGAA CTATTTTAAT GCAGCTATGA AAAGGGAAAA AAGTGCCCAG TTCTTGATTT CTTAGATACT GAAGAGGACG TAGCATTTCA TTTATCAAAAT ATAAAGGAAAA TTATTCACCA TTTTGAAGCT CACCCTAGAC TATGAAAATT ATATTCACTG CAGAGCAATT ACTTCTGTCA TTACCTGAAAG TGATCAGTAT CTATCTTCCT TGTCATAGCA TGCATCTCTC AAAAAGGCCT CCACTCCTTT CCCTCACATC TGTGGTCATC ATGATT

SEO ID NO:1118: (Length of Sequence = 379 Nucleotides)

GACAGCAGCE TETCCAGGGC GECTGTGGAG GTGTTCGGGA AGCTGAAGGA CCTAAACTGC CCCTTCCTCG AGGGTCTGTA
TATCACAGAG CCAAAGACAA TTCAGGAACT GCTGTGCAGC CCCTCAGAGT ACCGCTTGGA GATCCTAGAG TGGATGTGTA
CCCGGGTCTG GCCCTCACTG CAGGACAGGT TCAGCTCACT GAAAGGGGTC CCAACAGAGG TGAAGATCCA AGAAATGACG
AAGCTGGGCC ACGAGCTGAT GCTGTGTGCG CCAGATGACC AGGAGCTCCT CAAGGGCTGT GCCTGCGGCC CAGAAGCAAG
CTACACTTCA TGGACCAGTT GCTCGATACC ATCCGGAGGC CTGACCATTG GGTGCTCCA

SEO ID NO:1119: (Length of Sequence = 233 Nucleotides)

CAATATTCAA GAGTCTITAT TGAAGACTIG AGATGGGACT TCCAACTCAG AGGATGIGGG AATCCCAGCT CAAATGATAC
AGGATAAACT GGGATGGGCT AGGATGGACA GGCTGTGGAT ATGGGAGTCA TGGGTCAAAG TCTTATCCCA GATGGCTCCA
GGTACAGTGG GCTTCCTGGG CTGGAAGCTG GGTCCTCCCC ACTTCATTCT GCTCAAAGCT TCTTGAAGGA GCT

SEO ID NO:1120: (Length of Sequence = 325 Nucleotides)

GAAAAAACAA CCATACCCIT NCTTTIGAGG AAAACITACA AACITTATAA AGAATAAACA TGAATCINCT TAGAAAGITC CAAGATAACA TACACAACIG ANICACCICI TCATATATAG GCACCACACA CATAAAGATG TAGCCTAAAT CACAATCACT TCTCACCAGG GATGGAGATA GGAATTTACA TTCTTGACTT CATTAAGTCT CTAATTTGGC AAAAACCTCC AAGCCTTTTA TACACATGCT GCGIGTAGGC CAGATCTCAC TCATTCTTAT AATTGTGCAA ATAATATGGA GACCAAAAAGG GCAGGGTTTT CATTT

SEO ID NO:1121: (Length of Sequence = 161 Nucleotides)

ATTAGTATTT TYSTCTGTAT GTCCTAGCIC TGTTCAACAA CAAATTTTINC TAGTTCTYST TIATTTTIQAT TTGTTATACA ATGGAAGCAC AATGTTATAA GGAAAGGTAA TTTTAAGCTA ACAACCAGTG CACAGCCTCA GGTTTTAAAT TACAACCACA SEO ID NO:1122: (Length of Sequence = 181 Nucleotides)

CATCITITIA CATCAAAGIA CTACCAAGIA AAGAATITAA AAATTACITG TCTAGTCATG ATATATITIC CINCIGCTGC
TGAAAAATCC CTGTCTTATT ATTTCATGIN CCTTTATCAT TCATTTGATG ACACTGACAG CAACTTGCTG AACAAGITTA
AGAATAGCTG ATATTTACTG A

SEO ID NO:1123: (Length of Sequence = 174 Nucleotides)

CCCTAGAGTT AAATTCACC CATGAAACAT CAGCCACATG TCATATCAAT TCAAGTGTGT AACATTGATA TAATCGGGTA CACCACAGCA GCACTGACAG AAACAGAAAT GATTCAGAGA AAGCCAATTA AAACAGCCAG GGGATAAAGC AGATCTGTAT GACATTAGCT TTTT

SEO ID NO:1124: (Length of Sequence = 232 Nucleotides)

CTTTTAGCAG AGACGGGTT TCACCATGTT GGCCAGGATG GTCTCTTGAC CTCGTGATCC ACCCGCCTCG GCCTCTCCAA GTGCTGAGAT TACAGGCATG AGCCACCGCG CCTGGCCCAG GGAAGGCATT TTTNAAGAAA TAATAGTTGA ATTGAGATCT GATAAAAGAA GTAGGAGCAA AATNGGGGGG GTGCAGTTTT CCAAGAAGAG AAGACAGTAC ATATAAAGGG CT

SEO ID NO:1125: (Length of Sequence = 233 Nucleotides)

GATACTATEG GITCAGIGAC ATAGAGACAC AATIGAATTA GCAATGAGCT TCACTCAGGA GCCAGAGAAT GGGITIMINI CTAAGAGATG TITTAAGIAA CATITAAATG GCACTGCTGA TTGATACCAG CATCAGGAAG CTGAGGACAA GAGCTCTCTG AGAAGGAAGT TGCCATATTA CAGAAGIGAG GTGACCAAGC ACTINTIGIA GGICTGTACA TITTAGACATT AAT

SEO ID NO:1126: (Length of Sequence = 258 Nucleotides)

TTTTTTTTT TCCTAGGGGC CGCAAGACGG CTAATTTATT ATAATTCCTC CGCCGCAGTT GCCCTCTGGC GCCA....TCGC
AGAACGGAGC GCCCGGGATG CAGGAGGAGA GCCTGCAGGG CTCCTTGGGT AGAACTGCAC TTCAGCAATA ATGGGAACGG
GGGCAGCGTT CCAGCCTCGG TTTCTATTTA TAATGGAGAC ATGGAAAAAA TACTGCTGGA CGCACAGCAT GAGTCTGGAC
GGATTAGCTC CAAGAGCTCT CACT

264

GTGTGAATAG GCAAGCACTT TGTTTGTTGT ACTAAGGAAC TCAAAATGAT AGGCTTTTTG TCACCATGTG CTTCCAGGNT
CTCTGTTGCA TGAGCAGAGA TAGAGGATCT TGCACAAACA ATTAAATGCT CTAGCCATAA GTAGTGCAAG TTTCCNTTGC
TTGAAATTTA CTGCTGATAG CCACTTGGNC ACACCTTACT TCCAGAGGCT AGGAAGTACA GTTTTCCCAC AGTCTAAGAA
TGAAAGAGNA TTAACCACAG TAATGCATAG CACTCATACC ATGGATGACT GGATAATTTT AAAAGAATGG GAATATGCAA
G 321

ACAGCTCAAT GACTIATCAC AAAGCAAAGC CCCAAGAAGT CACCACCCAG CTCCAGAAAT AACACATTGA AAAGCTAGAA AAATCTCAAA TTGACATCCT AACACCACAA CTAAAGGNTC TAGAGAACCA AGAGTAAACA AACCACAAAG CTAGCAGAAG ACAAGAAATA ACCAAGCTCA GAGCAGAACT GAAGGCAATA AAGACACAAA AAACCTITAA AAATAGTCAA TGAATCCAGG AGCTGTTTTT TTGAAAAA

SEO ID NO:1129: (Length of Sequence = 163 Nucleotides)

291

CAGIGGIACA GCAGCAGCAG ACACGCATCG CAGAGCIGGA GAAGACGICA GCTGAACACA AACACCAGCI GGCGGAGAGA AGCAAGACAT CCANCIGCIA AAGGCATACA TGCATGCAAT CCGCAGIGIC AACCCCAACC TTCAGAACCI GGAGGAGACA ATT

SEO ID NO:1130: (Length of Sequence = 382 Nucleotides)

THITITITI TITITITITI TITITITITI TITITACIGI TCAAACAGCA ATGITIAGIT GIACACACA TAAAGICIAG CAACAATIAC AGGNCCAGIT TGAGIGICIG TITIGCITIGIT TICAATIGGG AAAITTAACI GIAATGICAC CGIAAGAITIG GCIGGGACIG GIAACATITA AGAAACGGGI TGINCITIGCA TCCCCTAGGC GIGGGCCTCT TGCTCCATCA GGACITIGGIT GIAGATGAAT GGCCCACAAG TCACCAGCCT TIGAGCAAGI TGIGTCCAGG TGGAGACAGG AAGAGGGIGG GCAAAGGGGA ATTCTATAAA GACACAGIGI NIGGGGCAGT GGCAGICAAC ATTCCACAAAC ATTCATGCAT CT

SEO ID NO:1131: (Length of Sequence = 406 Nucleotides)

ATGCTAATTC AGGCTCCACA GATAGINCTG GTGATGGGGT TACATTTCCA TTTAAACCAG AATCCTGGAA GCCTACTGAT
ACTGAAGGTA AGAAGCAGTA TGACAGGGAG TTTCTNCTGG ACTTCCAGTT CATGCCTGCC TGTATACAAA AACCAGAGGG
CCTGCCTCCT ATCAGINATG TGGTTCTTGA CAAGATCAAC CAACCCAAAT TGCCAATGGG AACTCTGGNT CCTCGAATTT
TGCCTCGAGG ACCAGACTTT ACACCAGCCT TINCTGATTT TGGAAGGCAG ACACCTGGTG GAAGAGGGGT ACCTTTTTTG
AATGTTGGGT CACGAAGATC TCAACCTGGN CAAAGAAGAG AACCCAGAAA GATCATCACA GTTCTGTAAA AGAAGGTGTA
CACCTG

SEO ID NO:1132: (Length of Sequence = 400 Nucleotides)

ATTITIGGIT ACTICAGGCA GGAGGGIAGA CATAGCACTI ATCIGGATTG GATGIAGCCA CAGGATIAGA ATTGITGGGI CATAAAATAT GIACATGITC AGCITTAGIA GATCITGCCT AGAGTITAAA AAATTAAAAA TIAAAATATI TITIAAATTA CAATAAATIC AGCIAATTIT AATTITAGAT AATTITIATA ATGIAGTIGA TCITGGITTI AACCAGAGCA TGINGCIGGA TITINCICCC CAATCGAACA CAGTAGAGAG AGAAGGIGGC GGGTTCTTAG TGATACCATG CACTTTTTIT TAGAACTICA GIGCIGIATC CCITCATTTA CAATGIATGA TGAAAAATAC TAAAGAAGGG ATNGIGGIGG TGGIGAGGGA GGCAGGAGAG

SEO ID NO:1133: (Length of Sequence = 347 Nucleotides)

CCCAGGGCGC GCCATCCATG GACGAGCTCA TCCAGCAGAG CCAGTGGAAC CTCCAGCAGC AGGAGCAGCA CTTGCTGGCG
CTCAGACAGG AGCAAGTGAC AGCGGCCGTG GCCCACGCGG TGGAGCAGCA GATGCAGAAG CTTCTGGAGG AGACCCAGCT
AGACATGAAC GATTTTAACA ACCTCCTGCA GCCCATCATC GACACGTGCA CCAAGGACGC CATCTCGGCC GCGAAGAACT
GGWTGTTCAG CAATGCCAAG TCCCCGCCGC ACTGTGAGCT GATGGCCGGN CACCTCCGGA ACCGCATCAC GGCTNATGGG
GGCACACTTC GAGCTGCGGC TGCACCT

SEO ID NO:1134: (Length of Sequence = 389 Nucleotides)

GGTCCAGGCC TGCAAGACTT GCCTAGTGAG AAGATATAGG AATGGGAACC CAGGTAACAG TCTGGCCACT TINCCATAGG GCTGCTGCAG TATGCCCAGG GCCCGCTCCA GTCTCTAGTA GCCTCANATT TTCCAGTACC TGGAGTTATC ATCAGTGAAG CCTGTGAAAAC AGCAAAGATG GCAGCCTACC GCTCCCTTTG GAAGCTTTGC CCTAGGGAGG TATGAATGAN CTTNTTGCTG GCCCAAACAC ACCTGTAGGA GGTGGCTAGA GACCCCAGTT TGGAGCTTTT TGCAGCTTTT TGCAGGCTTTT TGCAGGCTT CCCCAATCA CCCCAATCA

SEO ID NO:1135: (Length of Sequence = 402 Nucleotides)

GCAGAGGCTT AAAGAGTGCT TATTCACTGA GCCTTGCCCT TNCTTACTCC TTCCTGGGAA CCCATTTGGC AACAAGTGAA
GAAACCTAGG CCAGCCTNCT TGAAGATGAG GGACCAACGG AGAGAGAGGC TCTGCTGTCC TAGCCCTCCC ACAGAATAAG
TAAGCCTAGC CAACACCACG TGGAGCAGAG ATGAACCATC TCAGTTGAGC CCAGCCCAAA TTGCTGACCA AAAGAATTGG
GAACAAATAA ATAATTATTG TTTTAAGCTA CTGTGTTTCT GGGTGGTTTT GTATATAATA GTAGCTACCT GATACATTGG
GATGACCCCA ATTACTTGAA CTTCTCTTAG GCCTGTTTTA TCACGTGCAA ATAGGGGATA ATTTTAGTAA TTTNGGGTTG
CT

SEO ID NO:1136: (Length of Sequence = 381 Nucleotides)

CAGGIGGGAG CCACCACGCC CAACCCAGAA CICTITITAT TITGCAAAAT TGAAATICTA CCCATTAAAT AGCAACTCIN
CTITICCCIT CTCCCCCAAG CCCITGGCAA CIGCITTTCC ATTICTATGA CAATGICTAC TCTAGATACC TCATAGAGGG
TGAATCATAC AGTATITGIC CITTITATGAC TGGCTTATTT CACTTAGCTG CTATATTATT AATACCAGCT TTCTGGGGAT
ATAATTCACA AACTGCAGAA TTGAATGGTT TINAGTCTAT TCACATCGGA TATGTTTTTG AAGAGACAGT AAAACCAATC
CTTTTTTCCT TAGGTTCTCA GACACACAC TGCTTCTTTA TCTGGCAAGT CCCGTTATAA A

SEO ID NO:1137: (Length of Sequence = 325 Nucleotides)

TATTITITET ATAGACAGGE TCITGITATE TIGCCCCGAC TGGTCTCGAA CCCCTAGTCT CAAGCCATCC CCCTGCCTTG
GCCTTCCATT CCTCTACTTT ATACCACGGT TATTCACCAA GCTTGTCTTT GTTCAGTGTA CTTCCTCATG GAAAAACTGA
GGTGATATTT ACCCTGGTTT TTCTACCAGT GTGTAACTGT CGCTAGTACC AGCTCAAAAA ATAAGAAATG AATAAATGAG
TGATGACTAT CACTATGTTG CTCAGGCTGG ACTTGAACCC CTGGGTTCCA GTGATCCTCC CGCCTCAGCC TTCCAAGTAG
CTGGG

SEO ID NO:1138: (Length of Sequence = 422 Nucleotides)

CAACACACAT TAGCCTTAAC AACAAAGAGC TAATCTTATG TAAAGAACTC TTACAATTCA GAAAGAAAAA GATCCTAGTG
AAAATGTGGG CAAGAGATAG CAAAAAACCA GCCATATGAT AATAATAGTC AATAAGTGAA TCTGAATGAT GTTATCTNCT
TTTGTCATTT TAGAAATACA AATAAAAATG ATGATGAATG CNCTTGCTTA CTAAATTAGC AAAANCTGGG AAAAGATGAT
GATATTCAGG GTCAGATAAA GGGAAAAGGG TGCCCTTCTA TTGCAGTTTG GAAAGTAAAT TGGCACTGAC TTTTAGTGGG
GATAGTCTTG TAATATGGGT CAAANGTCTT CAAATCGTGT CCACATTTTG GGGCCTGCAA TTCCACTTCT AGGGATTTAT
TCTAAGGAAG TACCTAAAAA AT

SEO ID NO:1139: (Length of Sequence = 367 Nucleotides)

ATACCGAGAA GCATGCAAGC GGTGGCTCCA CCGTCCACAT CCATCCCCAA GCTGCTCCTG TTGTCTGCAG ACACGTTTTG
GATACACTCA TTCAATTGGC CAAGGTATTT CCCAGCCACT TCACACAGCA GCGGACCAAA GAAACAAACT GTGAGAGTNA
TCGGGAAAAGG GGCAATAAGG CCTGTAGCCC ATGCTCCTCA CAGTCCTCCA GCAGTGGCAT TTGCACAGAC TTCTGGGACT
TATTGGTAAA ACTGGACAAC ATGANTGTNA GCCGGAAAGG CAAAGAACTC CGTGGAAGTC AGTGCCAGTG ANGCGCTGGC
GGTGAGGGGG TAAACCTTTT NCATACAGCC TTCGAGGCCT CTCCACT

SEO ID NO:1140: (Length of Sequence = 412 Nucleotides)

ATCCAAAGGA TATAGGCAAG CATCAGATAC AGCCAAAGCA TICTITICCT AAAAGAGTCT GAACGCATCT NAIGCAACAC CCAAAAGTAT CCCTTINCTC CTCGTTACAG TATGTTTTCC CTTTGGAATA NATCATTACT TATTGAACAN TATATGGAACA AATATCTTAC AAAAGGAAGT CATTTCCATT TICTAACATC TTTTACATTG CACTAATTAC ATGGTTTAAA TGACTATCCC TAATCTCAT CCAACTACAC CCCATGAATT TNAGGTTTAT TTAATCAACC TAGTTAGACC AGATATATCC TTCTAAAATC

ATTIGIAGAT AGAGGATICI CCITTITIGCI AGIAAATACC ATTAACATAT TINCAGANGG CCIGGICIAG GGICATTIAT TCCAGGGCCI CI

SEO ID NO:1141: (Length of Sequence = 410 Nucleotides)

GTTAACCTGT GGCGCGCTCC GGGTATCCGG CGCCTGANGT TTTAGCTGCG GTGGCGGCG CAGTCGGGAC CGACTNAAGA
TGTCATTTGT CAGAGTGAAC CGCTGTGGTC CCCGANTTGG TGTAAGAAAG ACACCGAAAG TAAAGAAGAA GAAAACTTCA
GTGAAACAAG AATGGGATAA TACCGTGACT GATCTAACCG TTCATCGGGC AACTCCTGAA GATCTGGTAC GCCGTCATGA
AATACACAAA TCGAAGAATA GAGCATTAGT ACACTGGGAA CTCCAAGAAA AAGCTTTGAA GAGAAAATGG AGGAAGCAGA
AACCAGNAAC TTTAAATCTT GAGAAAAGAA GATTNGTCTA TCATGAAGGA GGNTTCTTTC TGATCAATAC CAGATGCAAA
GATGTGTTGG

SEO ID NO:1142: (Length of Sequence = 392 Nucleotides)

TTTTTTTTTT TTTTCCNGGG ATTGAATGTC TTTATTAAAT AAACGAGTAA ATGGTAGCAC AAATCACCAT CAATATTTTT GGAAGGATTG GGGACAAGAT GTCGAGTCAG AATTAAATTN TCCATTTCAG GGTCTCAATG TAGCTGAAGA ACTGTGCCCCA CTGATCAGTA TTACGTATTG CAAATGCAGG AGGTAAGGCT AAAATAGGAC TTATGCCGTT CAGAAGATTG AMTTTGAAAC CTTAAAAACT ATCATAATAG TAGGAATGCA TGTTAAGATT TGATAACTTT CTTTAGCTAG AGTTTTCAAC CCACAGTTAG GAGCAAAGTT GTAAAGTGAG TAGGTNTGAA GAAGGACAC TCTTTTGAGA AAAGAAATTN GC

SEQ ID NO:1143: (Length of Sequence = 200 Nucleotides)

ACTICCICIC TCCTGGCATC TGCTATAAAA ATAAGAAGGA GCAAATATTC TTGCCTCTTT TTATCACCTG ANCTGAAAAC CCATTGTAAC TGCCATGAAA ATAAGCACTG GTCCATGAGA CCAATGCCCA GAAAATTCAG GCTAAGATTC CTGGAAAGTG GGCTGTGGGC ATTATTTAAA ACACACACAC AAAATTTACC

SEO ID NO:1144: (Length of Sequence = 333 Nucleotides)

AACAGAAGCA TGITATTCA TTCCCATTCC CAGAAGGGA GTTAATGAAG ATAAAAATTT ATTTTTTAAG GTCTTTATTG
AGAGAAACTT TGITTTCTGA TATGAACTAT TGCAGATGIT TTTATAAATA CTTTCATTAA AATGATGIAA ACAGTAGTAC
CCAACACTGT AAACTCAGTG AAAATAGTAA ATGATTCTTT TATTACTAAG ACTGTCATGC ATTCTGAAGC AGFTGGCTTT
TTTTTTAACCA TAGGAAGTCA TTTCCCTCTA GCTCCTTCCC TTCTACTCTC CTGCTCAGAC CATTAGTAGG TACTTTGTTA
AATAAAAAAC TAG

SEO ID NO:1145: (Length of Sequence = 225 Nucleotides)

TGGGTTTCTG ATCCGAGAAA AATTGAAAGA CAAACATGGC TGGGGGAAGC AAAACGCTGA CACACAATTC AGGTGGCCCA GCAGTGCTGA CCTGCAATCC ACCCCACCCC AAGGCAGCCC TTTCAATCCA AAGTGGACAG AGTGGGCCTT ATCCCAGANT CACTCAGGAA GCTTCTTCAA ACATATGACT GCCACACCCG CCCCCAAGGT TCAGAAACAT CTTCG

SEO ID NO:1146: (Length of Sequence = 223 Nucleotides)

AAGGNACAAT ATTATTCTAA ATAATTTAGA TITGGAAGAC ATCAATGACT TITGGAGATGA TGGGTCCTTG TATATTACTA
AGGTTACCAC AACTCACGNT GGCAATTACA CCTGCTATCC AGATGGCTAT NAACAAGNCT ATCAGACTCA CATCTNCCAA
CTGANTGTCC CTCCAGTCAT CCGGGTGTCAT CCAGAGAGTC ACCCTAGAGA GCCTGGGGTA ACT

SEO ID NO:1147: (Length of Sequence = 389 Nucleotides)

ATTICAGIGG CCATTAAGAC CCIGAAAGIT GGCTACACAG AAAAGCAGAG GAGAGACTTC CIGGGAGAAG CAAGCATTAT
GGGACAGITT GACCACCCCA ATATCATTCG ACTGGAAGGA GITGITACCA AAAGTAAGCC AGTTATGATT GINACAGAAT
ACATGGAGAA TGGTTCCTTG GATAGTTTCC TACGTAAACA CGATGCCCAG TITACTGTCA TTCAGCTAGT GGGGATGCTT
CGAGGGATAG CATCTGGCAT GAAGTACCTG TCAGACATGG GCTATGTTCA CCGAGACCTC GCTGCTCGGA ACATCTTGAT
CAACAGTAAC TTGGTGTGTA AGGTTTCTAA TTTCGGACTT TCGCGTGTCC TGGAGGATGA CCCAGAAGC

SEO ID NO:1148: (Length of Sequence = 386 Nucleotides)

ATTAATTECT TECCATCATE AGCAGAAGCA AGCGTGACAA CAATTTINAT AGTGTAGAGA TTGGAGATTC TACATTCACA
GTCCTGAAAC GNTATCAGAA TTTAAAACCT ATAGGCTCAG GAGCTCAAGG AATAGTATGC GCAGTTNATG ATGCCATTCT
TGAAAGAAAT GTTGCAATCA AGAAGCTAAG CCGACCATTT CAGAATCAGA CTCATGCCAA GCGGGCCTAC AGAAGACTAG
TTCTTATGAA ATGINITAAT CACAAAAATA TAATTGGCCT TTTGAATGIT TTCACACCAC AGAAATCCCT AGAAGANTTT
CAAGATGITT ACATAGTCAT GGAGCTCATG GATGCAAATC TTTGCCANGT GTTCAGATGG GCCTAG

SEO ID NO:1149: (Length of Sequence = 364 Nucleotides)

GGCAACAGGG TGAGACTOCA CCTCAAAAAA TAAAAAAAAA GAAAGATATT ATTCAAGAAA AGAACTTAGG AGCCAGGTGC AGTGGCTCAT GTCTATTATG CCAGTACTTT GGC.AGGCCAA GGCAGTAGGN TCACTTGAGG CCGGAGTTC AGAGACCAGT CTGGGAAAACG TAGCAAGACC TCGTCTCTAC AAAAAAAAGTG TTTAACAAAT TAGCTCAGTA TGGTGGCACA TGCCTGTAGT CCCACCTACT CAGGAGGCAG AGGCAGAAGG ATGGTGCCAT CAAGGCTGCA GTGAACTAAG ATGGTGCCAT TGCACTCGNG GATGGTGAAC AGAGCAAGAC TCCATTGCCG CCAG

SEO ID NO:1150: (Length of Sequence = 267 Nucleotides)

GACAGGIGIA ATCIAAGCIT AAATAAACCC CCCGGAGGCI GCACAATINC TIGGCATCIC TCCCCTGCCC TCTCCATCCG
CATATICATI TIGGAGITIG GAGAAGIATC TAGAATCINC TCCCACCCCA AAATGCCCAG CAGAGCCCCC CCGCCGCCCC
CGCACCCCTT GGAGCIGCGG CTIGCTGAAT CGITGAGAIG TCIGANACIG TCGGGGITCC CIACCTAGIG CITCAACCAG
ATCACCTCAC TITTGAGITT CCTTCCT

SEO ID NO:1151: (Length of Sequence = 386 Nucleotides)

GGAAGACGAA GGAGGAGTAA AGGCATGINT CACATGGCAG CAGGCAAGAG AGCGTGTGCA GGGGAACTGC CCCTTATGAA ACCCTCAGAT CTCGTGAGAC TTATTCACTA CCATGAAAAC GGCACAGGGA AAACCTGCCC CTAAGCTTCA GTTACCCCCG ACAGGTCCCT CACATGACAC ATGGGGACTA TGGGGACTAT AATTCAAGAT GAGATTTGGG CAGGGACACA GCCAAACCAT ATCAGATACT TACCACATTA GACACTGACA GACAGCTCAC CACAGATTCT GGGCTCTATT CAAGGTGTTG ACTTTGATCT TTTTTCCAGTT GTAAATGTTT CATCCAAAAA AACTGTGATT TTGGCCATAAC TTTTTTCCAAG AGTTGC

SEQ ID NO:1152: (Length of Sequence = 239 Nucleotides)

GCAATCTITT GAGGACITA CITTGAGICT TIGTCACCIT TCCICIGATI TITTCACATG GITTAACICA GIGTACCCAA GAGTACTAGG TGCACTCAAT TCIGCIATTA ACTCIATAAG CAAGINCITA AGAAAGITAA TGITAAAAAA TAATCITAAA ATTGICITGA TAGGAAAAAT GIATTIGAAA TTAAAAAAAA TTCTTATGIT GACITCITGG TITTGAAACA ATGAATATA

SEO ID NO:1153: (Length of Sequence = 275 Nucleotides)

CAACCTCCTC TICAGTGTCA AAAAAGCCAC GGTTAGACCA GATTCCTGCC GCCAACCTTG ATGCAGATGA CCCTCTAACA
GATGTATGTT TIGTTTCCTC CTTTCATCTC TAATAATTGA TTTACCATGT TTTTCTAAAA TACTTGTTAT GTCTTTNCTT
TAAGAAGTGA CATATATTTA TGTTTAGTTA CTGTTATTCA AATATAGCCC TGACCTCAGT GCTAAACTTT ATAGTTGATT
TTAAAATCAA AAGTATTATT TTGTGGGACT TTAAG

SEO ID NO:1154: (Length of Sequence = 203 Nucleotides)

CCTAAATCTT AAACCTTACA ACAGITAAAT AAGACCCCIT TCAAAGGGAT TAACACACTG AATATTATAT ACATACAGAT TTATATTTAT GCGCTATACA CATATATGGN CITTATCTGT ATATAAATAT GTGATGATAA TGATAAAAGG ATAATGATTA CACGTAGGAT AAACATTTAT CAAAAATTGT ACTATAAATA ATA

SEO ID NO:1155: (Length of Sequence = 343 Nucleotides)

GAAAACAAAA CACTAAGCTA TTTTGGAACA ACTGITCTAC ACAGAAGAGA GCTTCTCTTA ATTTAAAAAA AAAAAAAATC CCAAATAGGC ATTTTAGGC ATTAACCAAA AAAGAGAATC CAAATGAAAT ATTATACTTG ATGITCAATT TTAATAGCAT CTTGATAAAAG GIATGCTTCC TTTCATTTGA NTACATTTCT GNACATGIAT GITATAAAAT CCAGGNAACA GCCAAACCAC AAGTTAACTC TTAACAATGA ATATACATAG TTAACCCTAT AGTAAGCAGC CCCTTTGAAA AGCACTGATG CACCCAACAN TTATATGGTT CCATTTCATA AGG

SEO ID NO:1156: (Length of Sequence = 396 Nucleotides)

COCACCAATT GCCATTAAAC CTCCCAATCT TTACTGGGAG GNTCTCTACT TACTGTTCA AGGCAAAAAG ATGATTAANC
TATCTCACAT GGITGTAATT TGGGCCTAAA ATAAATGACT CTAGTGGTAG CATTTCATGT AGGCAGGTCC AAGGAAGACA
GATTTGTAGA CAGAGTTGGG AAAAGGGTCA AAGAGCCAAT GAGTCTCCCT ATCCTGAGGG ATGCCTTGAC GGGACCACAG
CATGANCTCA TGTTTTCCTG AATCCATCTC AGTTCATGTG ACAGGATGGA AATGCTTCCT TTCTTAGCCA GTGTTGCTTG
TAACGAGTTC CCTGCAGCTC AGGGAAGGGA GCAACATGTA CTGCTTTGTT GCTTCCTGTA TAGAGAAGGC AGGAAT

SEO ID NO:1157: (Length of Sequence = 269 Nucleotides)

CAGGGICICA ATCOGICTCC CAGGCIGGAG TGCAATGGCA CAATCICAGC TCACTGCAAC CTCCACCICC CGGGITCAAG
TGATTCICCT GCCICAGCCT CCCIAGIAGC TGGGACCACA GGCACCICGCC ACCGCAACCA GCCAACTITT GIATTITGIAG
TAGAGACAGG GCTTCACCAC GCTGGCCAGG CTGGTCTCAA ACTCCTGACC TCAGGTGATC TGCCTGCCTC GGCCTCCCAA
AGTGCTGAGA TTCCGGCGTG AGCCACTTG

SEO ID NO:1158: (Length of Sequence = 190 Mucleotides)

CTTATTAGIT AATTCCACGG CAGATTITCA TITCIATCGA ATATATTATA TGTAGAAACT AGGCCCTTAA ATAATTAAGC TGACTTINCC TATTAGITAT TCCTTAAGAT AAAATTATGC TGGTGAAAAT NACTGINGAA TITCICAAGA AATTAAGCTC TATAGAGGCA TAAGTAATCG AAAGACTTIT

SEO ID NO:1159: (Length of Sequence = 340 Nucleotides)

GGGCACTGAC TTCCTGGGAG TGTAAGCNNC TCACCTGGAC CCCACAGCCA GTGAGCATTA GTGCTTATAT TCCATCCTCC
AAAGCTCTTT CTTCATACCA GACCACACAT GTGGCCCAAG GAGGGATATT TACTCTGCAC TTTTAGAGTT CTAGAAAACA
TTGTTTAATG GTCTGGCATC ATCTATATTT ACTTGGCATA CAGTATAATC CTAGTCCTCG ATGAAAAAGTT
TTTNATGAGT TAACCTTATG GGGTGATGGG ATTTATGGGA TTATTTCCAC CCTTAAAATG ATTTTGTGGG GAAAAAAAGTT
GTACTAATCC CTAATTTAGG

SEO ID NO:1160: (Length of Sequence = 215 Nucleotides)

GTAAACAAAT CAATTAACAT GATTATCCCA GACCITTCIT TTCTTACTGG AAAAAAGAGG GCATTAAACT GGATGATGAC
AATAACACCA TAACTACAAG CITTTATAAA AGTCCITTAT ATACAGTGIT AATACAGTGA AAGNICAACC TTATTGAAAG
AGGTCTGGCT TCTGCCCTCA GCTACTGGGA AACAATCACT AGGCCTCTGG CATGT

SEO ID NO:1161: (Length of Sequence = 298 Mucleotides)

AATCITIAAA ACTACTITGA ATCITATAGA AACATCAGAA TCITITGAAT TCAAAAGAAG CCAGGGACTC TAGCCAAAGT
GGAGTGGIIT TITAACTCAA GGATTTAGGA CCITGGCTGA ATACAAACAT TGAATGATTA CTCAGTAGGI GCCAAAGCTC
AGGACTTTAG ACAGAGTCAG AGTCCAGTIT GINCTGAAAC ACAATTTGAT TTCAACTATT GITTTAAGTG AGAGAGGAAA
GTGACATTAT TATGAGTGTA AATTINCTGC TITTTAAAGTA GAAGTTACTG ACAATTGA

SEO ID NO:1162: (Length of Sequence = 163 Nucleotides)

GAAATAAGAA ACAGCTIGIA TATAACTAAT GCTTIGAGGG AGAAATICAA ATGGCTATGA AAAAATATIT ATAATICAAT GATAATAAAA ATCTTACACG TTAAAACTIG AGAATGTAGT TAAAGCAATA CITGGNCATA ANCITAGCAC ATATTAGTAA AGA

SEO ID NO:1163: (Length of Sequence = 393 Nucleotides)

GCCAACACCA GGAGCATTIT ATTCAGATGI TAAATGAACC AGITCAAGAA GCTGGTGGTC AAGGAGGAGG AGGTGGAGGI GGCAGTGGAGGI GAGGTGGAGGI GAGGTGGAGGI GAGGTGGAGGI GAGGTGGAGGI GAGGTGGAGGI GAGGTGGAGGI GAGGTGAAGA GAGGTTAAGA AAGGTTAAAG GCATTAAGA TTCCTGGAAGG ACTTGGAGG ACTTGGAGA GGGACTTTIT TATATCTCAC ACTTCACACC AGTGCATTAC ACTAACTTGI TCACTGGATT GCCTGGATG ACTTGGGCTC ATATCCACAA TACTTGGTAT AAGGTAGTAG ATT

SEO ID NO:1164: (Length of Seque ce = 260 Nucleotides)

SEQ ID NO:1165: (Length of Sequence = 330 Nucleotides)

CATTGGTATT TAAAAATGAA TATTAATATA ATGAAATGCN TTTGCCTTTT TGTAGGCATA ATAAGCCAAA TACTTTTTTA
CCCAAAATAA TTTTNAGAGA AAATGATGTA ATGAAAAAATT GTACCATGAA TTAGGAGGCAT AGTTTTNCC ATTTAAACGT
CACCATTACT TAAAAGATGA TTGATTATTG CTATACCAAA TCAGATGAAC TCTGTTCATC ACTTTCCINC TCTGTCCCCA
AACAATTTGG TTCATTCAGA CTGAAATGTT TGTGTCTTCA ACTTATTAGA ATGGAAGATA ATGCAGATAT TTCTGTGGGA
AATAAAATAA

SEO ID NO:1166: (Length of Sequence = 312 Nucleotides)

ATTGGAGATG CCTTTGCCAA ATTTNCCCAT TTTAAATGGC CAGGAAAAAC AATAATTATT TICCTGATGC TGAGGTTTTA
TATCTTAGTA GAAGAACTTA AACTATGACT TGTATTCAAG TXTAACACAA ATTGTTTTAA ACAACCACTT TTCAAAAGCA
CCTGGGACAA GATCACTTTG AACATGACAC TATTATACAA AGTGTAATAT TTATTTTTAA ACAACCACTT TTCAAAAGCA

GITGTGCATA CATTCCAAAG AATAAAATGC TAGCTACTAG GTTTTGAGAA GCAGAATAAA ATATGATACT GA

SEO ID NO:1167: (Length of Sequence = 305 Nucleotides)

SEO ID NO:1168: (Length of Sequence = 342 Nucleotides)

AAGGITTIAG TGATGATICA GIGAGAAACA TATTIGAAGC AACAAGCACA GIAACIGGAA GCIGIAGGIA CICAATAAGI
GICAGITTCC TICCICITCI AAAAGCIGIG CITICAAGIC AATIGIATGI CIAGAGICGC ACIGICIGGI ACAGIGGCCA
GIACIAGCCA CATAIGGCIC TCAAGIACIT TAAAGAGGGC TAGICIGAAT TGATAIGIGI CATACATGIA AAATACITTA
AAGAGGGCIC ATCIGAATIG ATATAIGCCA TGCATGIAAA ATACAAATCA GATTICIAAA ACITIGIACC AAAAAATACC
ATAAAATAAC TTACIAATAA TI

SEO ID NO:1169: (Length of Sequence = 397 Nucleotides)

GAGACGGAGC TCCNICIGIC GCCAGGCTGG AGTGCAGTGG CACGATCTTG GTTCACTGCA AGCTTCACCT CCCAGGTTCA
CACCATTCTC CTGCCTCAGC CTCCCGAGTA GCTGGGACCA CAGGCGCCCA CCACCACGCC CAGCTAATTT TTTATATTTT
TAGTAGAGAC GGGGGTTTCA CCGTGTTAGC CAGGATGGTC TCGATTTCCT GACCTCGTGA TCCGCCCGCN TTGGTGTCCC
AAAGTGCTGG GATTACAGGC GTGAGCACCA ATGCCCCAGCC TTTGGAGACA CTTTTGATTG CCACAACTCA GGGTAGGGAG
GGCTGGGAAA TATTACTGGT GTGTAGTGCA TCGAGGCCCAG GGATGCTGCT AGACATCCTG CAATGCACAA GGACAGG

SEO ID NO:1170: (Length of Sequence = 422 Nucleotides)

GITTIAAAGC CICIGGACAG AGCAGIATIT CGITIAAAAC TITGITTITC TIAAAAGCIT ACAGIGITIG GCIAATICIC CICCCCITIT TACAAGACGG GGGCCGGAGG GIGGACACTG GIGGCAGGIT AAGGGATACT GICACITIAA GAAGCCIGCA GAITGAAGIG TAAACAIGGA GAAATTAGGG GCIGATITIT TAAACIGIGI GAGATATIAA CCAGCCGCCC TGITATAAAA TCAGGGAAATC CAAACAGCGA TITACACCGA TIAACACCCC CITTATATAT TITINACAAA AATACACTGA GAAAATAATC AAACGITTIC ATCICICTIG TCTITTITIG TTTITTAAAA GIGTCAAAAG TCTACATNIA AATATAAAAN AITAAAAGIT AAACTCTAGC CCTTCAGIGA GG

SEO ID NO:1171: (Length of Sequence = 384 Nucleotides)

TCTGAATGGG TTGGTGAAAG GTTACAGGAG CAGACAGCCT CCACACCCAG GCTGCTCTTG GCTATACAGG CTACCTCCAT
CCCTGANTGT TGTAATAGGA AAGTCTAAAC ACACAGAAGA GGAGCACAAA ACCAATAATT ATCACACATT CAAAATAAAA
CTAATCCATA AAGAAAAGTA CCAAACTCAA CAAAGACAGC AATGCCTGAA AACACTGGGC TGTATCAGCA AATAGAACAA
AGAAAAATAN GCATAATTAA AACAGTAGAA GGTGAAGGAT AATTTTTAAA ANTTAGATAT CATATTCTGA TTATTGAAAT
AAAAAACTTA GTAGAAAAGC TTAACTGAAG AGGATCAAAC CTGAGGAGGA CCCCGCCAGT TTTG

SEO ID NO:1172: (Length of Sequence = 410 Nucleotides)

GAGAGAAAAA AAAAAAATCT TTTAAAAGCT GCCATCTGAG GTGATGGCTT CTCTGTACTT ACGCCATACC CCAGANTACA ATAAATAAGC AATTAGAAAA CGTTCAAGTA TGAAGGGATT TCCTCCTCCC CGCCAAAAGC ACTGCTCTCT GAAGGAAGCT GGITTCICTG TAGCIACACC AGCIGITCAG AAAGCICATI GGACCIGGIT TIGAAAATAA AACAAAGITA AAACCCIGGG AGGAGITATT GINCAGIGIG GAGTACTCAG GCITTCITAT AAAGAAAAAA AAAGGITATC TGGTACCAAA GIGIGCACCT ACAGACCCIC AGGTACTGCC CIGIGACTIC NCIGIATGAC ATCACAAGGC TGCCAAGIGC TGCTTINCTA GACTAGGGAG TIGGTGAGGT TITGCTAG

SEQ ID NO:1173: (Length of Sequence = 274 Nucleotides)

GAGATCIAAA TGAAATTIAT AAGAAAATIG TGGGITCIGC CCAAGATGAC ATCIAAATIG AAGAAGGIAC ACAGTGAGIIT
TAAAGGATCA ACGAGAGAAA CIITIATIAT TCATTIGCAC AAGAAGACAC ATTCAGTATC TGGATTATCC AATATATGGA
ATACITIGAG TIGAAATGAT TAAAGGGIAA TCITIAATCA TIAATTAACA AATCATTAAT TAANCAAAAT AATATTIAGC
AAATTAAGCA AGINCTAAAG GCTACATGCA AACT

SEO ID NO:1174: (Length of Sequence = 326 Nucleotides)

AGAAATTAAA ACACTITAAT ATAAACATIT CCAGAATATA GACTGACCTT ATATCAGTAC TITTINGAGAC CGITTTAAAA CTATATATCA TCTAAGTITA TTATAGACTG TTTCATTTTC CACTTTCAGA ACTAGAAAAT GCAAAAATAC ACTGCAAATT AGATTTAACA AAGAAAAAAT CAGTTTAAGA TATTTCATAC ATATTCCTTG GNGAAAGCTG AGACACATAA ACACAGNAAA ACACAAATAA AATACCACCA ACACTAACAC AAAACCAAGG AAAGAACTGN TTTTGTAACG CTTGGTAATT CTGTCCTTTA AAATAA

SEO ID NO:1175: (Length of Sequence = 426 Nucleotides)

GCAGTCAGGA TGGACACATT AGAAAGAAAC ATTTTAGTIT CAATGTTACC ATAAAACCAG AACGAAAAGC AGCATGCTGT
ATTATATTIN NCAATTTAGG TTCCATTTCT AACTCCACCT AAAATGAATA TGAACAAACT CATTTTTAAG TGTTTGTCAG
TCAAATACAA TAATAGTCTA AGTTTATTCA CATATGTACC AACCAAAGCC CAATAAAGCT AAAAGGAAGC CAAGTGTAAT
AAAAAGGCAG CTATAAGGTC TTGTGTTTGA NTTTTTACCC AGCAAGAAAT AAATGATACT TAGTAATCCA TCTTTCCCCC
CCACTGCCAT CCCTGCACAC ATCTAAAATA GCCTAACTTC ACCTATTCTA ACTTCTGAAA TTGTTTTGGG ATTCCTGTTT
TACTTTCTCA GAGTGGATGG TATAGC

SEQ ID NO:1176: (Length of Sequence = 301 Nucleotides)

CTAATCCTCA ATCCTATCCC TTINCCTCTT AGCCATCCTC TCTAATTINT TTAACCTAAG CCTGTGTGTC CTCAGAAAAT AGGTTATGCT GITGGTGTG GTGGTTGGTA ATCTCCCTTT TTACTCCAATA CTATATTTAT AAGANCCNIT TAAGTGGTTG TATGCCTCTA CTTTATTGCT TCTGACTGCT GCATGGNATT CCCATACTCAT GTCCACCACA CTTACTCATT CTCCCTCTTG ATGGACGCTG AAGTTGCTTG G

SEO ID NO:1177: (Length of Sequence = 331 Nucleotides)

GCAATTCTCC TGCCTCANCT TCCTGAGTAG CTGGGATTAC AGGTGCCTGC ACCACGCCCG CCTAATTTTT GTATTTTTAG
TAAAGACAGG GTTTCACCAT GTTGGTCAGG CTGGTCTCGA ACTGCTGACC TCATGATCCG CCCGCCTCAG CCTCCCAAAG
TGTTGGGATT ACAGGCATGA GCCACCAAGC CCGGCAAATC CATGCTTTTA AACATTACTC TGTATGGTGT GATAATGAAC
AGTCACTGNT ATCTGACTGT TCATCTGTGT GGTCCATCTG TATTGAATAA AGGAGGAAGG AGTTGAAGAA TAAAGGGGAA
AATCTTGCAG A

SEO ID NO:1178: (Length of Sequence = 325 Micleotides)

GAAATINITG GAGAGAATAG TCATACCTAC TITAAAAGAG AATAAATTGC CTTTCCTAAA INCCTCTGCT TCGCTCCTTT CCTGGCGTTG CTCTGGAACC TTGTGAGTTA TATGTATGAT INCTGTACTC TGATATCCAT CAAAGTGCAT AACATAGTAC TCATGATGCA GTAAGTACAA ATCTTTTTTG AAAGAGATAT TGCTTGINAA CATTTTGGAT TTATAACATT GGCTTATAAT ATATACAACA TCTTTATAAA TGCCACCTCA GTTGGGTTTT AAGCCTTACA AGAGTGCTAT GAGTAAATAT CACCCACTTT AAAAA

SEO ID NO:1179: (Length of Sequence = 297 Nucleotides)

CCTTGGGAAT TGTTTCTTGG AAATTAAAGC ATGTGTCTCA CACAAACAGT AGAAGGCATT GAGCATTCAT TAGTCTTTCC TCAAGAAGAT ATCAAAATGA GACTAGAAAC TCTCTGGTGA ACACAGAATG CTCTGAGGGG GNCCAAGGTA CATTATGACC TTAAAAACGAA CTCCTTCTCC ACTGGCCCTA TTACTCACTG TGGAAAGCAC CATGCCAGGC ACAGCAAGAG ACTTAAGAAC ACCTACAAAG GAAGATCTCT NCCATCCACT TGTGTAATTA TCTTTAAAAA GTAATCC

SEO ID NO:1180: (Length of Sequence = 278 Nucleotides)

GCTGCTTGGG ACTTGAAATC TGTGGCCGAA GACCNGTCAC TACATAACTT CAAAAATAAT CAACCACCCT CCCTTCCCAA ACCACCCAAA TTCACTCATC CAGCGTTTAC TTTTTTGAAT CCACTCAGAA CTTTTTNCTG CGACCCCCCT CCCTAAATG AGTTGGGTGG GGGGAAATG AATACTGAGT TGGCCTTTAT TTTTTAAAAG ACTTTTTGAT CCAATGAGGC CCCCTAAATA ATTGAGTTTT GGGTCCTGGT TGGTTTGTTT TATTTTGT

SEO ID NO:1181: (Length of Sequence = 331 Nucleotides)

AATTGAGTTA CAGGAGAATA CTGTGAACAA TTGTACAGCT AAAAGTAATA ATCTAAATTA AATGTACACA TTCCTAGAAA CACACAAATC ACAAAANCTG ACTCAAGGAG AAATAGAATA TCTCAACAGA CCTATAACAA CTAAAGATAT GGAATCAGTA ACCAAAAAGCC TTCCAACAAA GAAAAGCCCN GGANTAGATG ATCTTCACTG ATGCNTTCTA CCAAACATTT AAGAAAGATT TAACACTAAT TCTACTCAAA CTCTTCCACA AAAAATATGA GANGAGTAGA GAAAACTTTC TAAAATATCT TATGAGGGCA GCATTACCGT G

SEO ID NO:1182: (Length of Sequence = 345 Nucleotides)

GIGIGINIAG AGGATGGAC AGGATGCTGI TIATTINCCC TITCITGGAA ATGGACCITC TGTCCCTTCC ATTTGGACAC CACAGTGGAA GCTGGTGGCC TGGAAGGAAG GATTAGGTCA TGGACATTTG AACAGGTGCC TTGGGCATGA TGTATAGATG CAGTCATATA TACCTTGCTG GGNTGGGGTG CCACCTCCAG TGGNCAGCTC CAGATCCAAG GAGCAGCCCC CTGGGGATGG ACCCCATTCA TTCATCATGA CTCCCAACAG TTTTTNATTG TGGAAGAAGA AACTTINGCA TTATAGAGAC ATCATCACAA AACAGTANAA ACAAAATCAA CCCTG

SEO ID NO:1183: (Length of Sequence = 272 Nucleotides)

ATGGAAGATT CAGAGATCAA AGGTGAAGTC CTCTATCTCG GCTACTACCA ATCAGCCTTC GACTGGNATG ATGAAACANC CAAGGCCTCC AAGCAGCATC GTCTTAAACG CTACCACAGC CAGACCTATG GCAATGGGTC CAAGTGCGAC CTTAATGGGA GGCCCCGGGA GGCCGAGGTT CGGTTCCTCT GTNACGAGGG TGCAGGTATC TNTGGGGGACT ACATCGATCG CTTGGACGAG CCCTTNTCCT GCTCTTATGT GCTGACCATT CG

SEO ID NO:1184: (Length of Sequence = 335 Nucleotides)

ACATTITICCA AACTCAGITIG ACTCACCTCA NATITIGCCAT TOCAATTACA GGGCCTCGAA AGAGTCAGCA CTCAGCCTTG
CTCAAGGNTC AGATTTAGGG GITGCCCCCC GNCCCCGCAA CCTCCCACCT ATTGITITCAA ATGICCTCAA GACAATCACC
ACTGITATTAA GAGAAAGAGG CATGGGGGCA GAGCAACAAG GAAATTAAATG AGGCTTTAGAACCT TAAACCACCC TTGGGNCCCA AATCTGCATG AGCAGGGGGT GGGCTATCAT GCTACAGANC CCCAAGGAGG
ACATTITITCC CAACA

SEO ID NO:1185: (Length of Sequence = 383 Nucleotides)

GAGAGGIGAG CAGGCGIGCG GGGGGGGGAC TICTGCAGAG AAAATATTIT TAAAGTCATA AAACCATGAA AATAACAACT
ACTGTACGIT TIATITIATA GAAATCAAGT AGTATCTAAT AGACAAGGGA AGACATTGAT CCATAAACTI TITAAAGAAA
ATTTGGTAAT CICTTAAAGT ATTTGTATGG CTITGAATGG GIGINCTITT CTAACTTTGT TITAATTTTT ATGATACACT
TATAATTGIT TCAAATAGGC ATTTGINCAT TITAAAACTA CTAGAAGTTA CACTGAAGAA AAGCATTCAA AAGAAGACTT
TTGGACAAAA AAAATTGITG AATGAGTGAA ATGCCTGAGG TAGCTCAATT TACCAAACAG GAA

SEO ID NO:1186: (Length of Sequence = 373 Nucleotides)

GGGGCTCAAG GTGTGCATGT NTGAGGGAAG AGAGAGAGAG AGAAGGCCGC CTCANAGGTG ACTITCAGCC TGCNAGCCTT
CTTCCCGGGG CGCCATAAAC GCCCCCAATT TCCCAGCTGC TAAAAGGAAGA GGAAGGTACC TGTNCGTGCA CGCAGACGGG
AAGGGCTGGG GAAGCGGGAG GACTGAGAAA AGCCAGATCT TAGCAAAGCA ATGTCTCAAG ATGGTGCTTC TCAGTTCCAA
GAAGTCATTC GGCAAGAGCT AGAATTATCT NTGAAGAAGG AACTAGAAAA AATACTCACC ACAGCATCAT CACATGAATT
TTGAGCACAN CAAAAAGGGC CTGGGTGGAT TTCGGAAGCT ATTTCATAGA TTT

SEO ID NO:1137: (Length of Sequence = 365 Nucleotides)

TCCACGCAAT ICTGAATAAA GTITATTAAA TAATATGTAC AGCAAATGTA GTAATTCAAC ACATCTATTI ATCAAATCAA
TCCACTGCAA TGAAGAAAAA TAAATGANCA GAAAAATCTA TGTCTGCATA GGNCATGCTC TCAGTGTGTA ATTTAAATGG
CAATACTTTA AATTAATTGG TTATATATAA TGTCAGTTAT TTTCCTTTCA GAATATAACC TTTTTTGTAG TAACCTATTC
TAGCAATAGG GCTTAATACG NCTGCAGATA AATAGGNCTG CAAAAACCAA AAACCCAAAA TAATGAAATT NAAAAGGGGA
AAAAAACTGT AACTGNGNIC AGAGTTACCT TTCCTCCCCC ATAGG

SEO ID NO:1188: (Length of Sequence = 350 Nucleotides)

AGAATGGCT TACATTTATT TTAAATTCA CTAAATACAA ATCITGATTG TCATGCCAGT TTTAGATCIT ATTAATTINC AGAATGGATA AATTCAAATA ATCATAAATT ACGGTAACTT TTTATTATAC CAAGGTGTTC TAATGCCATC ATATGANGAC AGATGCTTCA AACAACCTGC ATTAAATTAT ATTINNAATA AAATTAAAAT CTATTTTTAA CCTATTTGTA GTCACAAACC GAAAACGTGT CGNCTTTACC TTAGAGCTAA AGGCTTACTT TATGCATACG GGATATTTAA TAGTCTACAA ATCAAAGGTT TAAAACAGNCC CTTAAAAAATT CCATATATTC

SEO ID NO:1189: (Length of Sequence = 393 Nucleotides)

GCAAACTINC TCACTTCCTC AAAGAAGAGT AGTGCACTAA AAAGAAGGTT GCACCCGGAG AGCATGTAAA GTGTCTCAAG
GGGGACATCT GAAGINCCCC GTTCCCAGGG AGCCCACTGG CTCCTCACAA GTAATCTAAT GAAAGCTATG CATTCTCTCT
GGGCTCCTCA TATGAAAAAN CCCAATGTAT GANGCAAAGC CTAGAAAGGA TTCAATACTG GAGAAATGCA CACAGCTACC
GATAAAGACA GCTCAAAAGT CCTAAGGCTG CTGACATGAA CCAGATAATT GGTGGCTACA GTTGTGCCTG CTAAGATTTG
GGTGCATGGG GCTTCGCTTT GGTTAGCTCC CATGGTCTTC TTTTTCCAAA AAAAAAAAA AAGNCTTCAG GTT

SEO ID NO:1190: (Length of Sequence = 365 Nucleotides)

AGIGIAAACA TICUCCATATT TAATAGTACC TITAAAATAA GCATTACTAC ATTTAAAATG GITCCAAAAT GIATCIATAA AIGGIAATAT AAATTAAAAA ATACGAACIT AAAGIGAATA AATTTTTAAC CITAGCTATG GTATAAATAA TIGGIAAATGT ATAGIGIACC TNIGAGICAT TAAAATGICT TAAAAGATAA CAGCTTGTTA CCAGAACATT AGANACCATA GCCATGATTC TCAAGCENTA ACAATCTACA TTIGNIATIT NCTIGGCCAC TGCATTCTIC AAATGANTAA TAAATTTCCA GAATTCCCAT TCCCATGGIG TTITTCCCAA TAGANCTTIT TCACACTCGA TGTTG

SEO ID NO:1191: (Length of Sequence = 303 Nucleotides)

CCCGGAGAGC TGCCTTCCTC TTCTACCACG TGAGGACACT GCAGGAAGAC AGCTGTCTAA GAACTAGGAA GTGGGCCCTC
ACCAGACATT GAATCTGCGC TCCTTGAACT TGGACTTCCC AGCATCCAGA ACTGTGAGAA ATAAATTCAT GTTATTTATA
AACCAACCTG TCTATGGTAT TTNTTGTAGC AGCCTGCAGC TCTCTATCAC TCTTGTTTAT AAGAGGCTGA AGTTTACTTT
ACCTCAGGCA GAGCTAAGCA AAAAAGATTA CATCCCGATT ACAAGATGAA AGTAAACAGA ATT

SEQ ID NO:1192: (Length of Sequence = 315 Nucleotides)

SEO ID NO:1193: (Length of Sequence = 313 Nucleotides)

CEAATTAGTG AACTGTGCTT CAGGTTCAGG AACCTGGTCT TAGCTCCTTG CCTGCTGAGA TTTTGAGTTA CAAGTAGAAT
TCTCCAAAAAG CAAAACACGT AAAAGTCATT TINCCACTCT TTTGGTCAAG CACATGTAAG CTTTCAGGAC CAGGTGGTAT
GCCGTTNCTG AAAGTGAGAC ACATGCCCCA GGGAAAGGGT AATTTTAAAAA TTCTTCCCAT AGGTCCTCAT CCTGTTCCTC
TGCTATGTCC AGCATCCTTN AGTCCCAGCT GCAGGGCCTA TATTTAAATA CCCTCATGCT TTATCGCTTT TGT

SEO ID NO:1194: (Length of Sequence = 341 Nucleotides)

GATTTAAAAG CAAGTNATTT TNAAATCCAC GAAAGATGCC TACCITGGNT CCINCICIGG TCCITATTAG CCACACCTCT CTIGACAGGC AGAGGAGTTA GGAGTGAGGG GATATTCCCA CCAAGACCCT ACAAATTGCA CTCITAGGCC ATGCCCTGGG TACCCCAAACT CTAGAATTCC CTCCTCAAAG GGACCTTAAC CCAACTTCAG AGCCTATATA GGCCAATTCC TTGGTCCATT TTCCAAGGGG TGGNCAAAGG ACAACCATTT TNGGGAGGGN GANGGGAGTA GGATGAAGCT TTGGNCACGT GGGTCTTGGG CAAATCCCCAC ATATCCCGGA A

SEO ID NO:1195: (Length of Sequence = 239 Nucleotides)

TEATEGATE TETETEGAA ATGGAGTCTC GCTCTGINNC CCAGGCTGGA TEGCAATINC NOGATCTCAA CCCACTGCAA CCTCCGCCTC CGGGGTTCGA GCGATTCTCC TGCCTCANCC TCCTGAGTAG CTGGGACTAC AGGTGCGCGC CACCATGCCC AACTAATTTT GGTATTTTTA GAGACAGGGT TTCTCCATGT TGGTCAGGCT GTTCTCAAGC TCCCAACCTC AGGTGATCA

SEO ID NO:1196: (Length of Sequence = 291 Nucleotides)

CCATGCTTGG CTCAGGGCCT GGGGCGGGT CCTGGGTAGA GTCCTAGCCC CAGAGCCCCA GCCCCTCATG TCCTGCCGCC CCTCACTGAC CAGACGATGA TCGGCAAACCT CTTGAGAAAA CATGGCAAAG GATTAGAAAA GGGCAGGGTG AAATTNCCAA GCCACTCAGA CGGAACCCAG ATGATCTTCA ATGCAGCCAA GGAGCTGGGT CAGCTGTCCA AACTCAAGGT TCACATGGTA CGAGAAGAAG CCAAGAGCTT NACCCCAAAG CAGTGCGCGG TTGTTTGAGT T

SEC ID NO:1197: (Length of Sequence = 303 Nacleotides)

CTTCATATIT TTATAGCTGG GGTCAAAATA TGCAATTTAA AAATAAATAT ATCCATTINC CTATTCTIAC ATTTATGAAT ATAAAANTAA AATCTAAGAA ACATAATGCT GCCAACTAAT AGTAGTGGAG GAAAGGAAGC TGAGAGAAGG ATAAATATAT TANTITAATC ATTACTCAGA AAAGGCAGTA AAAGATACTA TCTATAGCAG GCATCAATAA ATATGANCCA TGAGCCAAAT CAGGCTTACC ACCTGATTTT NTAGGATAAA GTTCATTGNA AACACAGTTA CAGTGTCTTT CCA

SEO ID NO:1198: (Length of Sequence = 318 Nucleotides)

CTCAATTTCT TCTCATCTTT TINATGCTAT TATTGTCATA TAAGTTACAT TCCTATACAT TGTGTGTCCA ACACAAATTT
AAAATTATGC CATTGTCTCT TAAGTCATAG AACAAAAGAG ATACAAACAA AACATACATT TATCCTGTCT TTTATATTTG
CCTATGCAGT TACCTTTACC AGTGTTCCTT ATTTCTNCAT GTGGATCTGA GTTACTGTCT TTNAACTTCA ATCTAAAGNN
CTTTCAGTCT GAAAGACTGT AATTTNAATT TCTNGTAGGG GTAGGTTAAC TAATGATTAA TTCTCAGTAT TCTGAGGA

SEO ID NO:1199: (Length of Sequence = 326 Nucleotides)

TCTAGTTATT CTGAGAACTA CAACCAAGAA AAGAGGGAAG CACCGGGTTG GCCAAGGCCA TCCGGAGACT TGTCCTGCTG
GGTCATTTAA AAAGCTTTTC TAGGATAACG TTGGCTTTCC AAGTGGTTT CCAAGCTGAT GTCTTTCCCA CTGAGGAGAA
GCTGTAGGCC TGTGGACTGC CAGGTAGGAG GAGGTTGAGG TTTAGAGGAA AGAGGAGAGC AGGAATGGGT TGTTTNCAGT
GGGGCTGTTC CCATGGACTC ACCAAGAAGA AATCGAGGTG CTGATGGGGC TGCACAAGTG CTTATCAGAA ACAGCTGTAA
CAAGTT

SEO ID NO:1200: (Length of Sequence = 341 Nucleotides)

GGGTGACAGA GTGAGACTCA GTCTCGCTGA AAAAAACAAC AACATTGCTT TACAGTGTGA TTCCAGTTAC AGAGAATATT
CACATAGGTG CATAAATAAA TGAAAAAAATT ATTGGTTAAT GTCTCTGTAT GTTGGGATTC TCAGTGATTT TNTTTTNCTA
CTTTTNATTT TTNATAATTC CTCCAGTGTG TTGGTGTTAG CTTTATAGAT TATATCAAGT AACCTTTTGC TGCACCAAAA
AACCCCCCAA ATCTAGTGGA TTAAAACAAA ACCATCTTAC AATTTINNTC AGAACTGTCT AAGGCTGGAT ATTTTACTGG
GCTCTCTCCT GAATGTGGGG G

SEO ID NO:1201: (Length of Sequence = 312 Nucleotides)

GTCTTINITA CCCTGCTAGC AATAGCTCTC AGITTCAGAG GCACAGTCTT TGGAGACCAT TCAGCACTGA GAAAGCAATA
TTIAGAACCT ATTGCAAAAC TGGGCCTGAG TTAGGCATGG TGATGAATGC ATCAGCAAGG AATAGAAAGT NCTTATCGTG
AAACCCTTCA ACCTCAACTA TGCCTTCATA GACACACACG TTCATGCACA TGTAGGCACA TGTACCATCT CACATCTTTC
ACTTTCCCGA GATGCCATAT ACAATTACCT ACATTAATAN CTGTAGCACT ATACCTTTTT GAGCCCGAGA GA

SEO ID NO:1202: (Length of Sequence = 344 Nucleotides)

GGAAAATAGC CAGACTGGGT ATTATGCATG TAACAAATGA GGACATTGTG CATAAGAAAG GAAACATTAG TITTCTGTCA
TCCTGGGCCA AGTACCTCAT TACAGTAAAT GTGTGTCTTT GGAAACTCTT TGCTTGTNCT GATGGCGGTA AGCATGGGGT
CCCAGGCAGG TTCAAAGGCT GAACTGTAAG AAATGGGCAA GACAATACAT TTTGTTTTGG AAGGAATTTC TCATGGGATA
AGTTTCCCAA AGCTTGAATT ACAGGCTATG AAATAAAGCA AATAGATGGA GGAGAAAACA AGTATTGTTT TCAAAAAAGGT
ACCAAGTCAA TTCTATTTAA AGGA

SEO ID NO:1203: (Length of Sequence = 370 Nucleotides)

GITCITTATC TICCTCCTCT TATGIGCACT ATGIAATGIC CICATCATTT TAAAAGIGAG TIGCTATIGG GCGGGCGCGG
TGGCTCACGC CIGTAATCCC AGAACTITGG GAGGCCAAGG TITTIGGCTC ACTIGAGGIC AGGAGITCAA GACCAGCCTG
GTCAACATGG TGAAACCCAG TCTCTACAAA AAACACACAC AAAAAATTAG CTAGGCATGG TCGCACACAC CTGTAATCCC
AGCTACTCGG GAGGCTGAGG CACCAGAATT GCTTGAACCC AGGGAGGCGG AGGGTTNCAG TGAGCCCAAG ATCGTGCCAC
TGCACTCCAA GCTTTGGGGT GACCAGAANC GAGACTTTCT CAAAACAAAA

SEQ ID NO:1204: (Length of Sequence = 346 Nucleotides)

CTCTTTAGAA AGCCTGCCTT GGCTGGGCCT GGTGGCTCAC CTCTAATCCC AGCACTTTGG GAGGCCAAGG TGGGAGGATT
GCTTGAGCCC AGGAATTTNA GACTAGCTGG GGCAGTGTAG TGAGACTTTG TCTCTACCAG AAAAACCGGG CGTGGTGGCG
CATGCCTGTA GTCCCAGCTA CTTGGGAAGC TGAGGCAGGA GGGTTGCTT GAGCCCGGGA CGTGGAGGTG GCAGTAAGCT
GTAATTGTGC CACTGTACTC CAGNCTGGGT GATAGAGTGA GACCCTGTAT CAAAACAAAA CAAAAAAACAA AAACCTGCCT
TCTNGGGATT GGGCTTCTGG GTTTTT

SEO ID NO:1205: (Length of Sequence = 292 Nucleotides)

TACAACGAGA CACTIGAGCA CACGCGTACA CCCAGACATC TICGGGCTGC TATIGGATIG ACTITIGAGG TICTGTGTGG
GICGCCGTGG CIGCATGITT GANTCAGGTG GAGAAGCACT TCAACGCTGG ACGAAGTAAA GATTATTGTT GITATTTTTT
TTTTTCTCTC TCTCTCTCTC TTAAGAAAGG AAAATATCCC AAGGACTAAT CIGATCGGGT CTTCCTTCAT CAGGAACGAA
TGCAGGAATT TGGGAACTGA GCTGTGCAAG TCCTGAAGAA GGAGATTTGT TT

SEQ ID NO:1206: (Length of Sequence = 336 Nucleotides)

SEO ID NO:1207: (Length of Sequence = 319 Nucleotides)

TECCTCANCE TECAGAGTAA CIGGGATTAC AGGCGCCCGC CGCCACGCCT GGCTAATTTT TGTATTTTTA GTAGAGATGG
GATTTINCCA TGTIGGCCAG GCIGGTCTCC AACTCTTGAT CTCAGGTGAT CCACCTGCCA CAGCCTCCCA AAGTGCTGGG
ATTACAGGCA TGAGCCACTG CGCCTGCCTC CATTTCCTTT TTATAATTCA TCCCTGAACT CCCTTAAGGT AGAGAAGCTG
TTTGATCGTC CCAGCCCCTG GGAGGCTGAA AGGTAACTTN ACCAGCTCCA TGCCTGAGTT TAGCACCTGC TGTGCCAGG

SEO ID NO:1208: (Length of Sequence = 357 Nucleotides)

GAGATGITIA AAAATGAAGI GGAAGITITI TGITTITGIT TIGITTITGC AGAAAAAAGA TITTIAATGG CITGAATGIN
CTGCCATAGI TGCGICAGAT TGICAGAAAA TIATGITGIA CATCTGAGAG AGAAAAGAAG AGCCITTIGA GGAGCTGCGC
TAAAAATTATT TITTGITTAG TCICTIAACT CITTGGCTIG AATGAGTCAT TGACTTTCCT TGCCAAGATA GGGITAGCAT
TTGITTTGIG TITTAAAAGC AGGCCAAGGG ATTGCCACGA GGGAGACAA CCTGAGCAAC TGAAGGAAGG AATTTCTAGA
AAATGIGTIT ACCAGTTGIT TIAGICTGAA TGTGATT

SEO ID NO:1209: (Length of Sequence = 362 Nucleotides)

CCCATCTGCT CCACCCAAG AAATCAGACA AAGTAAAATT TATTGAGACA GACAGAAATG CACCTACTCA GGACTACAGT
TAAGCATTTA CTATTAACCA AAGAGTTGTG TTCACATTCC AGATAAGTCT ACGTGGAAAA GCATTCAGAA TTTACTAGGT
TTTTINCTACA TCACTATTTC ATCTACAATA GGGACAACAA ACTGACACTC AGGATTTGAT GGGCTCTCAT TACAATGCTA
TACATTTAAC AGGGNCAAAC ATCAGTGACT TTGAGGAAAA AGTTATAAAA NGACCAAAAC CACCCACTGT AGGATGGGCT
CTTGGATGTT ACTGTACAGC GTGGGTCAAG GTAACAACGA GG

SEO ID NO:1210: (Length of Sequence = 349 Nucleotides)

GAGAAGATAG TAGAGAAAGT CAGCGITACA CAAAGGAGAA CCAGGAGAGC TGCCTCTTTT GCCGCAGCTA CCACTTCCCC
TACTCCCAGA ACTACAAGAG GTCGTAGGAA GAGTGTAGAG CCACCTAAGC GTAAGAAGCG GGCCACAAAG GAGCCCAAAG
CACCAGTCCA GAAAGCTAAG TGTGAAGAGA AAGAGACTCT GACCTGTGAG AAGTGCCCCA GGGTATTTAA CACTCGCTGG
TACCTGGAGA AGCACATGAA CGTTACTCAT AGGCGCATGC AGATTTGTGA TAAATGTGGC AAGAAGTTTT TCCTGGGAAG
TGAGCTGTCC CTTCACCAGC AAACAGACT

SEO ID NO:1211: (Length of Sequence = 344 Nucleotides)

TTTTTTTTT TTTTTCAGG GAAGAGCTT ATTGCTTCCA TGGGGGGGC CTGGGACGGC TGCCACAGCT TGGGTAAGCT CCTTGGGCCT CANTICCCIT TGGTCAGGC TAAAGGCAGA ACCCAACCAC CTGGCAGINT TGTTGCTGAA ACCTAGAACA TGGGCAAGT TGGTGAGTCC GGGCCTGCGG TAGTCCTATG GNTCAGCTGC AGCTGTGGAG GGGAGCTCTT CCCAGCAGGC GGANTGGGCG TCACCCTCCT GAGCTTTAAA GTTCTTTCTG CTATAGCCCT GGGGCGGTCT TGTTGGCTCC GAAGGAATGG GCTCCAGGGT TTCCCCATGG GACA

SEO ID NO:1212: (Length of Sequence = 364 Nucleotides)

AAAGAAAACC TGGTATTTC ACCATCCTCT CTGAAAATAA ATACTTTGAC TTGCACTGAT TACTACTTCA TCAGCATTCA
ACTCCGCTCC GTGGCACTCT GTGTGAATAA TTTTAAAGGC AGATTAAGCA TTCTAAAAAT AAATTCTATT GGTAAATTAG
GATATCAGAT GCTTCCATTA TAAAAGCCTA TCCTATTCTG TACTCTCAGC TGGCAGTCAT ATCCAGATCT CAAGCTACTC
TGGCTCTTAT TGAACAAGAA CCTATTCCAG GGNGGAGGT TTTGAAGAGG GGATCTCTCA TGGTTAACTA GAGNCAGGAA
GAGGCAGAAT TGCCCACATA CTCTNGCAGG AGTTAAATAA CAAT

SEO ID NO:1213: (Length of Sequence = 302 Nucleotides)

CTAATTITIG TATTITTAGT AGAGACGGAG TICTACCATG TIGGCCAGGC TAGTITCAAA CICCIGACCI CGGATGATCC ACCCGCCTCG GCCTCCCAAA GIGTIGGGAT TATAGGCATG AGCCACTGIG CCCGGTTACT TITTCCTTT TTAAAACACT GAAATTGCTG TATCTACCAC ATTAACATT TATTTAAAAA AATTTGTTAA ATAGCATATG TATGTAAATT TAATATTAAT ATACCTCTTT TITTGTCCTT CTTTAGGIGG TIGGAGCCTA GGGATACTTA CTTACTGATT TT

SEQ ID NO:1214: (Length of Sequence = 317 Nucleotides)

CTAATTTINC AGACAGGITT ACATGIAAAA GGCTAGGTAT TTAGCCACCT CAGCATTGAT TAGITTIGGA TGICTAAGCT CTGITACACA TGGCTTCCCA TGGCTTCACT CTACAAAACA TATTINCAAC GTGAAGGNIA CATCTACAAG AAATCTACAT TTCAAGGGIT TTACAAATCA ATCTTGTATC TTTCCCCTGA ATTGACTCTC ACAGACCCCG TCCCCTTGIN ATTNCCTTTG CCCAGCTTAA CCGTCCAAAG TCTACTTAAAA TGCAGCTCAA AAATGTTAAG ATTGGGCAAC AGATTTACAG TTCCTGT

SEO ID NO:1215: (Length of Sequence = 276 Nucleotides)

ATAAGGTATT AAACAACTAT TCTTGTACTT GANITAAAAA AAAATCAAGC TGGGTGCAAT TGCTCATGGC TGTAATCCCA
ACACTTTGCT AGGGTTAAGT GAGAGGTTCA GCCCAGAAGT TCAGTACCAG CCTGGGCAAT ATAGTGAGAC CCCTTCTCTA
CAAAAAAAAAT GAAGAAATTA GCTGGGTATG GTTGCATGTN CTGTGGNCCC AGCTCCTCGG GAGGCTGAGG CTGGAGGNTC
ACTTGGGCCC AGAAGGTCAA GGCTACAGTG AACCTT

SEQ ID NO:1216: (Length of Sequence = 354 Nucleotides)

GCATAGGCAG CCCCIGCTCT TGCATTIACC TCCCACGIGA ACTAGCTCCT CAGICATTGC TCTGGAATAT GCAGTTGTGA TCTAGAAATT AAAGATGGGA TTAGGTAACC AGTGAGGTCC CTTCTACTGC CAGTGTATGA CTCTCTTCTT TGTAAATGTC ATATGTAGGG TTCTGTACAC AGGACATTTT CTTCATTGTA GTTCCTCAGA TGCATTGAGC TCTCCTGAAT GACTTAGCGG GGAAGCTCAG TTGCAGCTGA CCGTATTAAG GGTCCTCTCC CATTGTGCTG TGCCCGCTCG TTAGCGTAGG ATTCNTGCCC CACGGCCCTT CCTGTTTTCT AAGGGCTTGG CTTT

SEQ ID NO:1217: (Length of Sequence = 272 Nucleotides)

CTTCCCAGCT TTTGCTTGTT GTAAACAGCT GGCAGTGGTT ACATCTATAT TTGTTAAGAG GCAGAGCACT GTATTTTGTG
TAAGATAAGG TGCTAGTCTT GGCCAGGCTG CCAAGCTGGG GCINTTTAAA ATAAAAGTTT TAAAGAAAAA TTATAGCATA
ATAAATTACA CAATTTTATT GGAAAACTGA AGGTGTTCAA CCAATGCTAG TTTTTAAATA TATTTAGAAA TACTATTTCA
GGAAAATTTTA ACTACACTCA TTAGTCTTAT GG

SEO ID NO:1218: (Length of Sequence = 281 Mucleotides)

GTTGCCCAGG CTGCAGTGCA CITGTGCAAA CGCGGCTCAC TGCAGCCCCA ATCTCCCACT CTTAAGCAAT CCTCCCACCT CAGCCTCCTG AATAGCTGGG ATTACAGGTG TGCACTGCCA CACCCAGCTA ATTNCTTTAA TTTTGTTTTAT TTTTAGTAGA GATGGAGTTT CGCTATGTTG TAAAGGCTGG TCTGGAACTG CTGGCCTCAA GCGATCCTCC CGCCTTGGCC TCTCAAACTG CTGGGGGTTAC AGACGTGAGC CACCATGCCT GGGCCTGCTC A

SEO ID NO:1219: (Length of Sequence = 231 Nucleotides)

GTCTTCTCTC CCTCCTTCCC TTTATTGGCA CTGCCCGGAA CCAGGCAGCC AGCAGGGGAT GGGATCAGGA TGCAGTTGTC ATGGAAACGG TTGGGGATCC ACAGGAACGA CATTCATACA GGGACATTIN TGAAAGCAAA GCAAGAATGA NIGCTTTCCC GATCTCAGAC TGGCTGGATT CAGATCATTG TTTTGGCTGG TTCTCATTTT AAGGGGTAAG CAGTTTGCTA T

SEO ID NO:1220: (Length of Sequence = 409 Nucleotides)

AGICACTCAG AAACTTACTT TGCTTACAGC CTCATTATIG TITTITGIAT TIGITAAGAT ATTCCGGGG ATGACATATT
TTGCCTTAAA TTINCIAATT TTCCTGGCCA TTGCTTTCCT GTGATTCGAA AATGITACGG TAAGIGCTTA GTTTGGAAAC
TATACTGTCA ACATATATIG CATTACTTCA GCAGAGCTGT AGITCCATAA CATAATAAAA TGATGCTTTT TTTAATAAGA
AGATCATACA CATTTCATTA TGCCCTAAAA GATGAACATT CAAAGITCAC TTTTCTCTTG TTTTGATATG ACGGATATAT
ATCAGIAAAA TAAAAAATGC TGCAGNACCA ATATGCACTA ACTCAAACAT GCTGTGGATT TGTAGGGGCA CTGAGGTAGC
AATGTCAAG

SEO ID NO:1221: (Length of Sequence = 396 Nucleotides)

ATCIGAGATA CITIGICCIC ATGAATAAAT TAGITAGIAG AATCIAATIT CIAGATCCIT CATAATGGIA ATIGAGGGIA
AAAAATAATA ATGIAGIAGI CAATTITAGC CCITITAAACC TATGGGGAAC TGIATGAATA ACIGITIGAA ACIGCAGGGI
AATCCIGICA CACITGCAAA CACATAGAAG CAACAAGACI ATTICCTCIC ACACTITIAA TIAAAATAGI GCCIGAGIAG
ACITCCAGGG TAAGGIICAG AAATTINCIT TCIAATTICC CIGITITAAT GACCACTACT TITAAAGCIA TGCIGGGAAT
TCACITICAC ATATATCIAA CITACAGGAA ATTITIGAAG AGCCIAAATG TCIATGGGIA GATTCAATGT TICCIT

SEO ID NO:1222: (Length of Sequence = 350 Mucleotides)

GIATTINITI CIGGIACIC TICAIGGCCI GCIAGAGAAC TITACIAAAT TATAGICCAG TAGCIGGACA GAGCIGCAIG
TGIATICICI AAGICCACCI GIGCIGCIGG TCAAGAITAT TITGCAGIGI TIGGIGGIGI TGAAGAGGGAA TACIGITGIG
AAGGCIGAGI CAACIGCAIG ACAAINCICA TGGCICACIG GCWAAYGAGI TGIGGCAIGA CIAGAAAACCI CIGCIIGIAT
TCCCAGAIGA CAAGICACAC CIGAACAGCI GGATACIACI CGCAICCAAI TIGCIICCAA GITAACAIAT TINCAGAAAA
TATIIGGAIT TGGAGIACAI ACAAATAITT

SEO ID NO:1223: (Length of Sequence = 370 Nucleotides)

ATAAGCATAT GANTITATCT ATAGGCCAAG TITAATGACAT AACTACAAAG AAATGACTIG TITCACATGT TITAAACCAG
TGTTTTGGCT ATACTAACTT AGTGAGACAT ATTCTAAAGA AAAATAGAGA CGCAAAGAAG ATCTTACACT TITAATAGTCA
ATTTTGTAGT TGTAATATTA CTATCGATCA TTTTGTAACT CTCCTATATA GGGTGTAGGA TGGTGGAAAT AAGTAATTTT
MITAATGTTG TTAGGAACCA AGGCTATCAG TGTAAAATGA AGGAGTTACA AGCATAAGAT TGANAGACGG TAAGTAAAAA
GCTCATTAGT ATAGTTCCAA GTTTAACTTG TCAGGGATGA GCTCATGATT

SEO ID NO:1224: (Length of Sequence = 188 Nucleotides)

ACATGACCNA GGCCTGACCA AATCAGACTA AATCCTANTA CCTATACCAG AGTTATTGAG AAAGATAAGN TITGGCCTGC
NGGCCTTTGA CAGTGAAAGG NINTAGGCTT TGGAGCTCCT CAGGGCCACT GCTTCAGGGA ACCTTGCTGA CAGTGAAGCC
AACACAGATG AAAGCAAGGC CAAACATT

SEO ID NO:1225: (Length of Sequence = 353 Nucleotides)

CCCCAGCCAA GGGAGGCAGT NAGINAGTGT GGTACCCAGC GTGGGAAACC GTGCTTTTTIN CCATGGNACT NTGCAACCCA
CGGATTAGAA GATCCCACTC AGGAACCCAC GNCACTGGNA CCTAGAATGC CAACCCCAGA GCTGCACAGA TTCTAAACAA
CCTCTCANCT GGAATCTGCC TAACCCTGCA GAGCTCCTGC GGGGAGGGGT GACCAGTGCC ACANCTGCTG CTGCCTGCTG
CCTAAGCCAT TTAA

CAAAAAGITA G.IAAAACATG TAAACGTAAG TNATGAGGTA TITCATAGAT ACAGTGCCCA TACAAATNCT CITTCCCACA
ATTITCAACT GCCAGATCTC TIGCTTTAGT CITTTNCCT TATATITGGA GAAACAGAAG AGITTGACAT AAAAGTCCCT
TTGAGGATGT GAGGGTTGCA GTAGTTTACA GCAGGGTCAG AAAATGAAAG TAATAAAGCA ATATTTACAT GITTTTGTAT
AAGACCAAAA ATATTTCCTT AAAAAGTTGT TAAAAGTTTT TTAGTCCTAT AAACACTCAC TITTATAGGG CACATGATTG
TCTGTGTGAC TTCTCTTTCC AGAGGAGGC TITT

SEO ID NO:1227: (Length of Sequence = 352 Nucleotides)

GECATCIETT TITTIETTE TITTEAGATA GAGICICACT CIGICGCCAG GCIGGAGIGC AGIGGCGIGA TCTCGGCTCA CIGCAATCIT TGCCICCCG GTTCAAGCGA TTCTCCIGCC TCAGCCICCC AAGIAGCIGG GAGGIGIGCA CGCCACCACA CCCCGGTAAT TITIGIATIT TITGIAGAGA TGGGGTTTCA CCATATIGGC AAGGATGGIC TCAATTTCCT GCGCTIGIGA ATCCGCCCGC CTCAGCCICC CCAAGIGCIG GGATTCCAGG CGIGACCACG GCGCCCGGCC GGNATCIGIA GATTITAAAA GGCCCCCAGIG GTTCINATGC ACACCCCCAG AG

SEO ID NO:1228: (Length of Sequence = 387 Nucleotides)

AGITITICCAA GATIGAGIGA CACIATIGIA ATGAGAATCI TCACIGGAGC ATCAGAAGAA CIGATITICAA GCCAGITITIG
TIGGICAGCA CGGICAAAAC TICAGAAGAA TCITGIGCIC TGAGGCTITC CAAAGCTITIG TICCCCAGGG CAGTAACAGC
TICCAGIGIT GGCAGAGICI TIAGIATIAT CACCAGGGCA GCIGCACTGI GGCCTGIAGC CATCITICIC TITITAGIACG
ATCCCACCIG TCAGACITCI TGAATITIGCA CITCAAATTA GAGCCACAAT CAAATTATCA GICACGNIGI TTATITITIGI
CACCAGAGAA AGGACAGAGI CIGITICAGC AGAGTITIGGA GCCAGGTACT GATCTCTCIT CAGCAGG

SEO ID NO:1229: (Length of Sequence = 366 Nucleotides)

CTGATAAGGA GGTAATTICA TAGGAGCTGC TAAGATGGGC ATGAGGNICA AACTGCAAAG CACCAACCAC CCCAACAACC
TGCTGAAGGA ACTCAACAAG TGCCGGCTCT CAGAGACCAT GTGCGACGTC ACCATTGTGG TGGGGAGCCG CTCCTTCCCG
GCCCACAAGG CTGTGCTGGC CTGTGCAGCT GGCTACTTCC AGAACCTCTT CCTGAATACT GGGCTTGATG CTGCCAGGAC
CTATGTGGTG GACTTCATCA CCCCTGCCAA CTTTNAGAAG GTTCTGAGCT TTGTCTACAC TTCAGAACTC TTCACAGACC
TGATCAATGT TGGGGTCATC TACGAGGTAG CTGAGCGTCT GGGTAT

SEO ID NO:1230: (Length of Sequence = 343 Nucleotides)

AGIGGAGAGA AGCCCTATGA ATGTTTTGAG TGTGGGAAAT CGTTTTGCTG GAGCACAAAC CTCATTCGAC ATGCCATTAT CCACACTGGA GAGAAGCCCT ATAAATGTAG TGAATGTGGA AAGGCCTTCA GTCGCAGCTC GTCCCTCACT CAGCATCAAA GGGATGCATAC TGGGAAAAAAT CCCATCAGTG TAACAGATGT GGGAAGACCT TTTACAAGTG GACAAACCTC AGTTACCCTT CGAGAACTTN TTTTAGGGAA GGACTTTTTG AATGTAACCA CTGAGGCAAA TATTTTTCCA GAGGNAACAT CTTCCTCTGC ATCTGATCAA CCATACCAAA GAG

SEO ID NO:1231: (Length of Sequence = 406 Nucleotides)

CTCTCGCCGG GCAGCTTGGA GAAGGCGCAA TACTCTCCAG CTCCACCGTT ACTTCAGCAT GGCTGGGGAG GCCTTGGAAA ACTTATAATC ATGGTGGAAG AGGAAGCAAA CATGTCCTTC TTCACATGAC GGCAGGAAGG AGAAGTGCTG AGCAAAGGGA GGAAAGCCCC TTATAAAAACC ATTAGATCTT GTGAGAACTC ACTATCATGA GAACAGCATG AAGGTAACCG CCCCATGATT AANTTACCTC CCATGGGCTC CCTCCCGCAA GACGTGGAGA TTATGGAAAC TACAACTCAA GATGAGATTT NGGTGGGGAC ATTAGGCAAAC CATATCAATG TACATGTGTC TTTATGGTAG AATGATTTAT ATTACTTTAG GTATATAGCC AGTATTGGGA ATTGCT

SEO ID NO:1232: (Length of Sequence = 380 Nucleotides)

AGACCATCAA AGGCCCAGAG GAGAGACTCT TGGGACAAAT AAATATTTAA AAGCAGTTGC CTATGAGAAA ATGGAAAAAG CCACAAGCAA AGGTAAGATC CATGCTCCAA AAAGGCCTGA GAAAATCTTA AACCTTCTCC TCAGATTGAT CCCAAAAGCTT AGAACGAATA CCAAGATAAT AGCAAAAATC CTCCCTGGAA AAGAGTCAGT CTGCAAAAAAC CGGAAAAAGGA GGTTGTTTTT TCCACAATGC CTAATTTCTA ACAACAACAA CAAAAACTCA GAAAACATGG CCCAATAAGT GGAAGAAAAT AAAGTGACGG AAACCTTCCC CGGAGGAAAC ATAAGCTTCA GGCAAACTAG ACAGATTTTA GACTGTCTAA

SEO ID NO:1233: (Length of Sequence = 357 Nucleotides)

TTCAAAGTTT ATCACAACCA CCACCATCAA GACAGCAAAC CAAAGGGGCA TGGTAAAAGA AAGTTCCAGT GACTCTGGAT
TTGGTTCTAA TTTTAATGCA ACTTCTTGAT TGAGTGCAGG GTCAGCACTA CTTCGAAGTG GCTTTGGCGT TTCANCGGTG
GGTAATGGAG ACATTGCCAA ATTTATATTC TGTAATTTTN CGTTGGGTGA GGGGAGCATT ACATCATTAT ATAATGGTAC
TTCCTCAAGT TGCTGGTCAT CAGTTTCTGT GTCGTTGCTG CCAAAATCTA AAGATATGAT TGINTCTCCA GCGGCTGGGG
CCAGCAAAGT TAAAGCATCA GGTTCCTTCT TAAGTTT

SEO ID NO:1234: (Length of Sequence = 313 Nucleotides)

CCAAGAAATC TIAATINCTI TATTGTTTGA CITTTIGACI CAACAATTTI TITAAAACTI TITGTTTTI NCIGAAACGI TCITGTTGTT ATGAGCCTIT TGTTTTGINC TCGTTAAATG CACTCGACCC AAAATTGGTT TGGCATATCG AAAAGGAGAC CAAGGAGGGA GGGCCTGGGG CGTGGGGAGGT GGGGAGGAC; CCCGAATGGA CAGAAAGTTG AGGATAAGAG AAGAGGAACA TAGAGACACG CAGAAAGACA TGGGGAAAGA GTGTTGGAGA CCAGAAAGT GGAAGGCAAG CCAAAGCCA AAG

SEO ID NO:1235: (Length of Sequence = 386 Nucleotides)

CTCTCTCAGC ACAGCCTGGG GAGGGGGTCA TTGINCTCCT CGTCCATCAG GGATCTCAGA GGCTCAGAGA CTGCAAGCTG
CTTGCCCAAG TCACACAGCT AGTGAAGACC AGAGCAGTTT CATCTGGTTG TGACTCTAAG CTCAGTGGTC TCTCCACTAC
CCCACACCAG CCTTGGTGCC ACCAAAAAGTG CTCCCCAAAA GGAAGGAGAA TGGCAGCCTC CACATCTCGG GTTCAAGTGA
TTATCCTGCC TCAGCCTCCA AGTAGCTGGG ATTGCAGGTG TGCACCACCA TGCCTGGGTA ATTTTTGTAT TTTTAGTAGA
CACGGTTTCA CCATGTTGGC CAGGCTGGTC TGGAACTTCT GAGTGTAAAT GATCTGCCCA CCTTTG

SEO ID NO:1236: (Length of Sequence = 401 Nucleotides)

SEO ID NO:1237: (Length of Sequence = 372 Nucleotides)

SEO ID NO:1238: (Length of Sequence = 304 Nucleotides)

GGACAAAATT CCAATTATTG TAAATGTAAA AGAAAAGACA ACAAAAATAA GCTAGAAAGA TGAAAGCTAA AAATTCTATT
TGAACTATGT AAGATGATGA CAGATATTAA ACAGTAATTA GTCATGAAAC AATCATTTAA ATGCTTTINC CAGGGGAACT
GCAGAAGTTG AGACCCTCAA AGAGCATGCA AGCTAGTAGG GAGGCTGCGA CTCATACCTT TGAATCTTTC TGTTCTGCAA
ATTCTCAACT CTTACCAATT TAACTCTGCA GTACTGCTAT GGAAATTACA TAAGAGTAAA TTGG

SEO ID NO:1239: (Length of Sequence = 389 Nucleotides)

TGITATAACT GGCACTITAA TITGITTITG GAACTAGAAT TIAGGGGCAG TIGGATGAAA TIGCAAATIT AGAAGGGGAA
TAAGAATITC CIAGIGCTAT ATAAAGAAAT GATGATGGAG ACAAAAGCCT TGCTITCCTC TITITAGAAT TITATITINCGA
TITINAGCAT ACIGIGGGGC TITITAGAGCT AATATGATCT AAATINCAGAA AATITAATIT TCATAGIAGG CCAGGIGIGA
ATTACTIATG TITGCTATAG AATGCTTATT TAGACTAACA ATAAATITAC TITGCTTTCT AAGGCCAGIC AGCGAATGIG
GGGATGAGGC AGGATGTTIT AAATGAGCCA GAGATGATCC NCAAGGGGAA CAGTCGACAC AGAGGICTT

SEO ID NO:1240: (Length of Sequence = 365 Nucleotides)

CTCCAGCCTG GGCGACAGAG CAAGACTCCG TCTCAAAAAA AAAAGCCTTC CTTGCCAGGT GAAAGCAAGA GTGGTATGGA
ACATTTATTT AAACATAAGA AGCAGAAGGT TCCTCCTCTT GCAAGTATGT TTTCTCTAAA TGTAGCATTT CCACTGGAGG
AGGTFFTCTG GGTGGATGFT TAATATGTGA GGATTGINCA GCCAGGCCAGA TAACCAGGCC TCTGCATATA CAGATACCCA
CAGCCCAGGA ATCTTGAGAA CTGAATGGCC CATAACAACC TCTGGCACTA TCGGAGCTGC AGGGAGGCTT GGCTGGGGCT
ACTCCAGTCT CAGGCCCCTG TTTTTAGCGG GAAGTCACAA GGAGG

309

SEO ID NO:1241: (Length of Sequence = 350 Nucleotides)

GGGGAAGGCGG TAGGGTCTGC NCTGTCTGTN AGGGGCTTGT GGCTTGGCGG GTGGGCTTTG CATGGTCTCG CCTCTTGAGT
CCAGCCCCGT CCTGATGGGG CAGACTTCTG TNCGTNCTGC TTCCTGGGTG ATGTCAATAC TGAATGAGAG GGCAAGAGAA
GGGGAAAGGG AACCGCCCAT ATGTNCTTCA CGTGCTGCAA GGGGGCTGTN TGGTTCCCAT GAAATGGTCA GCAGAGACTT
TGGGATGGGT ATGACTCGTG GGTCACAGGG TTGACTAGAC AGAATCTAAA GAAGGTGGGT GCTTAGCTNG GAAGTCTTCA
GTAGGAACGG ATCACTGTGA AGCTCTAGGG

SEO ID NO:1242: (Length of Sequence = 392 Nucleotides)

CTCTTACGAG TGAGGITAAG TATTGAACAG ATATTTAAAA GCTATAAGCT TTTAAACAGA ATAGGCATAT TGCTGATACC
AGTATTTGAC AACCGCCTTG TTTTTTCAGA TAAGAAAACT GAAGCACAGA GACCATAAGG CATCAGCCTA TGGTCATTCA
CTTCGTGGTA GTCAGGTCGG AGGTCACACC AAGGCCCTCT GGCTACTGAT AATCTCTGTA CTAGGCTGCT TTTCAGTAAA
CTCTTGAATG AATGAAAGAA AGAACACATA CTGTTGACTT TTGAACTTGA ATCTAAACAA AACCTATGTT GAACTTTAAG
TCTGTAATCT AAGAACTATC AAACTTAAAC TTGTTACAAA AGGNGGTGAT GAGCACAACC ACTTTCTTTT GG

SEO ID NO:1243: (Length of Sequence = 377 Nucleotides)

GIGGGGCAGG CETGAGGTAG GGGTGGGGTG GGGATGACAG TCAACACAGC TTGGACCAGA AGCCCATGGC GCCTGGNTCC
CTGGAAAGGC ACAGGGCACA GACGGATGCC GCCTTINITG CTGGACACCT CCTGCCACCA TCCACAGCTC CCCCGTCACT
CCACGTTCTT GTACTTGGTG AACAGGTTGT AAAGAACCCT CAGGGTGGAT TTNAGGTCCA AGTTAACCAC GTCTTCAGGA
CGAGCCTTGG GTTINITNAG GCCTCCGTCC AGCATCAGCT CAAAGGCGAA GGACACATIN TGGACCTTCT GATCGAAGCT
TTCCCGGAGTC AGGTAGAAGT GGTGGAGAGG AACAAAGTAG TCTTCCAGAA GGCCCAT

SEO ID NO:1244: (Length of Sequence = 312 Nucleotides)

ATTITINCAT CAATGITCAT CAAGGATATI GGICIAAAAT NCICITITIC AGITGGGICI CIGCCAGGCI TIGGIATCAG
GATGATGCTG GCCICATAAA ATGAGITAGG GAGGATTCCC TCITTINCTA TIGATTGGAA TAGIITCAGA AGGAATGGIA
CCAGCICCIC CITGIACCIC TGGIAGAATI CGGCTGTGAA TCCATCIGGI CCIGGACITI TTITITCGITG GIAAGCIATI
GATTATIGCC TCAATITCAG AGCCTGITGI AGGICTATIC AGAGATCAA CITCITCCTG TITITAGICTI GG

SEO ID NO:1245: (Length of Sequence = 320 Nucleotides)

GEAGATCGTIG CACATCCAGG CCGGCCAGTIG CGGCAACCAG ATCGGGGCCA AGTITCTGGGA AGTCATCAGT GATGAGCATG
GCATCGACCC CAGCGGCAAC TACGTGGGCG ACTCGGACTT GCAGCTGGAG CGGATCAGCG TCTACTACAA CGAGGCCTCT
TCTCACAAAGT ACGTGCCTCG AGCCATTCTG GTGGACCTGG AACCCGGAAC CATGGACAGT GTCCGCTCAG GGGCCTTTGG
ACATCTCTTC AGGCCTGACA ATTTCATCTT TGGTCAGAGT NGGGCCGGCA ACAACTGGGC CAAGGGTCAC TACACGGAGG

SEO ID NO:1246: (Length of Sequence = 275 Nucleotides)

TTTTTTTTTT TTTTTTTTT ATCTGACAGC AATAGATTTA TTAAGTATCC COGAAAATAT AAACACAAAC CAGTAAAAAA CAAAACCGTA AAACGTCAGG CCTGGAGCTG CAATAAGACA GAGACAGGAG CAGCTCACAC GTGGCCTAGG TGGGGAGGAC GAGGCCATAA ATACTGCAGG AGGGCGGCAA GGGAGCCCTA GGGCGAGGGG AAAGCAGGGT NTCGGCAGCG AGATGGCTCC GGGGGTTTAG ACACTGCTGG CTTCGGCCCC GGCCG

SEO ID NO:1247: (Length of Sequence = 384 Nucleotides)

GGTCTTGCCG GAGAAGTACC CCCCTCCAAC CGAACITTIG GACCIGCAGC CCTTGCCCGT NTCTGCTCTG AGAAACAGTG
CCTTTNAGAG TCTTTACCAA GATAAATTTC CTTTCTTCAA TCCCATCCAG ACCCAGGTGI TTAACACTGT ATACAACAGT
GACGACAACG TGTTTGTGGG GGCCCCCACG GGCACCGGGA AGACTATTTG TGCAGAGTTT GCCATCCTGC GAATGCTNGC
TGCAGAGCTC GGAGGGNCGC TGTGTGTACA TCACCCCCAT GGAGGCCCTG GCAGAGCAAG GTATACATGG ACTGGTACGA
GAAGTTNCAG GACAGGNTCA ACAAGAAGGT GGTACTNCTG GACAGNCGAG ACCAGCACAG ACCT

SEQ ID NO:1248: (Length of Sequence = 225 Nucleotides)

AATTIGGAGA AGATAGAAGI TIGAAGTEGA AAACTEGAAG ACAGAAGCAC GGGAAGGCGA AGAAAAGAAT AGAGAAGATA GGGAAATTAG AAGATAAAAA CATACTITTA GAAGAAAAAA GATAAATTTA AACCTGAAAA GTAGGAAGCA GAAGAAAAAA GACAAGCTAG GAAACAAAAA GCTAAGGGCA AAATGTACAA ACTTAGAAGA AAATTIGGAAG ATAGA

SEO ID NO:1249: (Length of Sequence = 393 Nucleotides)

CATCTATAGT CCATACATAT CTATAATGGA CAGAAATATG AGAATGAATA AGCAAAGATA CTTATGTACA CCAATAATAA AGTAAGAAAG GTAAAAAAAA TCATGTAATA AGAAAAAAATA ACAACCCAGA AATTTAAGAN TTAAGTAGTA GTCAAATCTA ATTGGAATAA CTCACCTATA TAAAANACAA GAGGAAGGAA ACTTTATACA TAGGTCTGGA AAATATCACA ACTATGTTCC CAGAAGANTG TTTATCTCCA CAGCATCCAA CCTAGTGTCA TGCACACAGT TGGGACTCAG CCACTGTTGC CTGATTGATT ATGAAGNCAG TCACTGTGAT CAACCCCAACA GTAATTGAAC GTTCATTTTT AATANGGTCA GTGTTAAATC TGT

SEO ID NO:1250: (Length of Sequence = 391 Nucleotides)

CGTATGTATC TTINATITAC ACTGCACACC TIGCAGCATC CITACCITGC AGAGTACIGA GICCIGGCIT CATGAATTIN
ATGTCAAGTA AATGGGTTIT AGTCATCCCT AGTTCATGIG CATGINCCGA GAAAAAGGGG AGCTTCTAAA ACATGIGCGC
AAACCACAGG AAACAGIGCA ATCCTGIGIG TCTCCTATTC CACTTACTCC TCAAGGCCCC AAGGTAGGAC GCATGITTGG
TGGCTTTCTG GCTTACAAGT TCCAGIGCCT ACTCCCATTC CCTCAGAGGT TTGCTGTGAT CACTGAGGGG AAGCAGAATG
GAGCATCGTG TGGTCCTTAC TGGAGGACTC CTTGCAGCAC CTGAAACAAC CCAATGTTGT TAGAGGCAAA T

SEQ ID NO:1251: (Length of Sequence = 320 Nucleotides)

GCCTCANAAG GTCCTTCCCA GGCTTCTCGC AAAGGAAGGC ACTGCCTCIN CACACCTTGT GAAACCTTTC CAGGACCTCC CAGTCAGAGG CCGTCTGGGTT CTCACTGTCT GCAGAGGCGCC CTACAGCCTG TCTGTGGGTG AGCGTGTCTG TNAACTCTTG TCCATCTCTT CTGTGATCTG TGTGCTCCTC GAAATAACTG ATTTTNTCTC ATACACCTTG GAATCCTGAG TCCACAGAAC AGAGGCCTCAT ACAAAGGAAG CTTTCAAAGA GTGCTCATCG ATTTCTAGGN TTCTTGAAGA CAGGCACCAN GTTTTGTTCT

SEQ ID NO:1252: (Length of Sequence = 367 Nucleotides)

CAAAAAAACA AAACCAGTTA TGCAAAAACA AGAGTACAAA ATGCCCCTTT CTGAAGCTCA GTTTGAGAAA CTGATTTCGN
ATCTAGCTTA TTGATTATAC TCAGTTTCAA TTCTCCCTGT GCAAATAATA CATAAAGTCA TTAATGATGA TTTGATGANC
TGAAATCATC TTCGCTTAGG ATCGTTTGAC ATCATAACCC AAATATAAAA AAGTTATTCA AGATTCACAG AGATAAAACA
FTCCCTCCGA AACATAATTC ACCCATGTAT ATATAATANT TTTNGAACAT ACTTTTTAAA CATAAAATCA CAGTCAAGGC
AGTGATAGCA TTGCATACTC AGTGCATTAT TTCATGTAGT GCCTTCC

SEO ID NO:1253: (Length of Sequence = 393 Nucleotides)

TIGCTITCAA GACAACACTC AGTIGCTAAA CCCATTICCT TITCTITAGG ATATITICAT TGICTCCGAA TITTAGAGCT GAAAAGTGCC TIAGAGATCA TCTAGTITCAA CCTCTCCGTT CAAATGGAGA ACCTGAGCCA CTAAGNITCA CAGGNGAGTA AGATAATTGA GCAAACAACT CCAAGTAATG ACAGAAAATT ATAGGAGAAT CAGTACAAAC TGTGAGAATT TACTATGTTG TTAGCATCCT AAGTATGAGT TTAGAAAAGG TAGAAGTTAT AAGAAAAGTT AAATTGTTTT AATATGAATG GGATTCCACT GTTACCTTCA NGNIAAAATG GAGACATACT TTTTNCTTTA GGTATTATAG TTAAACGAAT ATTGTATCCN GTG

SEO ID NO:1254: (Length of Sequence = 377 Nucleotides)

CAAAAGCAAG GAGATGAGIT GAAAGCAGI TITNCTITAA GICATCAGITA TGGGATGICA GCAGAACAAA AATTAAAAAG AITAAATITNC CITTIGATCI AAAACITCCI TAGIITIGAGC AGIAGGIGCT ACAAAATITA TIACATATCI TAGIATCATA GITAAATGIA AIGIGITTAG GAGAGGAAAA CAAAAGATAC AITINCTITA AATTCATTAA GAAATTITCA AATTCACITT GIAGCCCATG CIGNATAGAA TIGGGCTGIG TIGGIACATI TGAAACACTG TITATGITGC TIGAAACACT TATTINTITIA ATCGCCCGATG TGATGATGCC TATGGCCCGAG ATCANATATA GCTAGATTGG CTAGGCT

SEO ID NO:1255: (Length of Sequence = 307 Nucleotides)

ACAAATGITA GCITTCICTG GCCTAGAAAA AGAATAGGNI CATCAAGICA TAAAACGAAG TAIGINATII CAGCACCICC ACAAAATGGC TICATCAAAG AAGAGAATCC CATCACATGI TACCICTCCT CICIAGGIIC TICAGCIGGG GCITIGCCIG CCCCTCTACC TAITGGCAGAA CCCACTGACT CGIGGNCTII CCAGCACTIC CACTTGCCIC CATTAGACAC TIAACCCCGC TGNCCGCTGC CICATGCCAG GGAGGGCCAA TCTCCAGNCA ATGCINCTGC TGGCTGTATG ATGACTG

SEQ ID NO:1256: (Length of Sequence = 326 Nucleotides)

TTGAGAAAAC TGCAGAAGCT GGAAGGTCAA TCTCTGACCT TCTTTCCTGA GACACCTTCA TGTGACAGGT GTCCCACTTT
ATGCCTGGAG GGAAGGAATG ATAACACAAA GATACCAAGA AGAATGTGAA GAGACCTTC TCAGTTCCCC CCAGTTCAAG
ACCATTATAT CGTACCCACT TTTGTCTAAT CANGCTTCTA TATGACTATC CATTCTTTAT CAAAACTAAA CATAGAAATA
TACGATTATC TCAATTTCTG TCTTTGNTTC TGAAGGCTCC TGTGTCACAT AAAACTTACA TTAAATAAAT TTGTATGTCT
CTCTTG

SEO ID NO:1257: (Length of Sequence = 224 Nucleotides)

TTTTTINAGA GGGATTCTCA CACAGTCACC CAGGCTGGGG TNCAGTGGCG TNATCTTGGT TCACTGCAAC CCCTGCCTNC NGGTTTCAAG CGATTCTCCT GCCTCAACCT CTCGAGTAGC TGGGACTACA GGCACCTGCC ACCATGCCCA GCTGATTTTC CTGTTTTNAG TAGAGACGTT GGCCAGGCTG GTCTCTTAAC TCCTGACCTC AGGTGATCTG CCCG

SEO ID NO:1258: (Length of Sequence = 329 Nucleotides)

CAGAGGITTC TITCCCTACC CITTGIGAAA ACCAATCAAT TACTAGATGA GIGGATGGAT GCAGAAAAAT CIGGCIGAG CCAAAGICCC TITTGGAAAT ACAAGCCATA ACATTCGAAG GACATCAGCG ACCITGGCIT GITTAGGIGA TITTINCITCC AGCIGCAGGI AGICTIGACA AGGAGCGITT AANCAGAAGG CICCAAGATGC ATTCCTTGIG TAGGIGGGIG AGAGCACTIC TAATGITTAAG TGGGGTACAG NICAGCIGCC CCCCCACGIA GCCIGGACAT CGICTINICC CCATAATCCT TNNCATCCCT ACAAGGICC

SEO ID NO:1259: (Length of Sequence = 374 Nucleotides)

GGTCATATGT TACATGCATG TTTGINCAAT ATGIGTATGT CAGGNCCATC TTCACAAATT INCATAGCCC CTTCTGTGAT CTGTTAAATA GGTATATTTA GCCAACCCTC TCAGCATAAA GCTCCTACCC CAGCTGCTCC CCCTTCCAAG TGCCTGCATC TECTCTTGGC TGGGAGCTCG CTTCCCAGCC TGTAGGATGG CCACCTTGAA GGCTGTAACC CTTTAGAAGA AATAAAGTCT CCTTTTCTAA ATTTATAGAT TGTATGATTG TTTTAAGCTA ACAATAGCAA TGGCATTATC ACCTCACTTT CTGTGTGTG GCTTAGCATA GTACCTGACA CATGGCACTT GAGTTGGTAG CTATTTTTTA ATAT

SEQ ID NO:1260: (Length of Sequence = 353 Nucleotides)

CTCAGTCAAA AATAGCAGCT GCTGAATTAG CATGGGCATA CCAGGCAAAT AAGCCTGCAT TGTCATAGCG TTCCCTTGAT
TGCNCTATGA AACTGAGTAA AGTTTCATTT CCTGATTCAA GAATTGCAGC TAAAATATCC TCTGGACAAA GAAGAAGGGA
AATTTTTTGA TAACAGATGT GTTGACTCCT TACAGTATAA AGCCAATTTC TGTCATATCT CACCAACAAT CCTGGTTTCT
ACAGTACATC AATTTTAAGT AATGGGCAA ATCATGGCAG CAAAAATATG TTCCCTCTAG CTGTTAGGGA CTTTGACTTG
NAAAACAGGN GTTTCAAATC ATCTTCTTCA TTT

SEO ID NO:1261: (Length of Sequence = 294 Nucleotides)

TTAAAACAGA CAGCTAAGAT TATAGGAATA TTTTAAATAA ACAGCATTTA TTTTAGACAC ATTTCAAATA GAAGCCACAA TAATCAAATA GATATTATCT GAAAAACGITT CAAAAATATT AACCCTTTAA ATGITCTTCT CTGAAAAAATT AGTTTATCTT TAACCAAATTA TTCTGAATTA TTGTGTCAAC ATATAAGGIT ATGCATATAT ATNCACTTGC TGGTCTCTAT GTTAAAGCAA ACTAGGTAAA AACTAGAGGA AATATCTGGA NCATAAAATG GTTAACAATT TACG

SEO ID NO:1262: (Length of Sequence = 292 Nucleotides)

ATGATGAAGG GITGGAGTGA TGCACCTAGA AGTGAAGGAA TGCCAATGGI TGCCAGCAAA GCACCAGAAA CTAGGGAGAA ACAAGGAAGG ATTCINCCAC AGTITCAGAG GGAGAATGGC CCTGCCAACA CTGTGATTIT GGACTTCTGG CCTCCAAAAC TATGAGACAA TAAATNCCTG TTGTCTTAGA CCACCCAGIT TGTGGAATTT TNITACAGCA GAACTAGGNA ACAAATACAG TTTTTTTTTTG CAGTAAAGAA GITTTAAATC TGGGTTATGI CCAATGTATC AA

SEO ID NO:1263: (Length of Sequence = 303 Nucleotides)

GGTTGAGGTT GTGGGTAGGA TGAGAAGACG ACAGGATGAA TCTTACCCCC CAGCTTTAGT GGAATTCTGT GAAACACCTG
GGAATGTGTT AGCATCAGGA GAATTCCTCT AAGGTATGAA GAATGACAAC CTGGGACCTT TCTTGTAGGT GGCTCTGAAC
CTAACTATTC CCCAAAGATT CCCAAGTGGT AGGAAGGAGG GGGTGCAGAG GGATATTAAT CATGGTCATT AAGTCTCAAA
ACATTTCTAC TTCAAGTGAA TACATTAACC ATGCTGAGGC AGTTGAACAA CTGAATGCGT AGT

SEO ID NO:1264: (Length of Sequence = 313 Nucleotides)

GGGACTACAT CAAGCACCTG OGOGACATCT GOGAGGGCTA OGTOCGGCAG TGCCGCAAGC GOGCAGACAT GITCAGCGAG GAGCAGCTGC GTACCATCTT CGGGAACATC GAGGACATCT ACCGCTGCCA GAAGGCCTTC GTGAAGGCCC TGGAGCAGAG GTTCAACCGC GAGCGCCCAC ACCTGAGCGA GCTGGGTGCC TGCTTNCTGG AGCATCAAGC CGACTTNCAG ATCTACTCGG AGTACTGCAA TAACCACCCC AACGCCTGCN TNGAGCTCTC CCGGCTTACC AAGCTCAGCA AGTACGTGTA CTT

SEO ID NO:1265: (Length of Sequence = 290 Nucleotides)

TTICTATGIG TAAGAGAAAA TAGAGATGGG TATACATACT GTIGTTTTT TIGAGCCGAG AAACIGIGIG ACCGGGGCCT CAGGIGGIGG GCATTGGGG CTCCTCTTGC AGATGCCCAT TGGCATCACC GGIGCAGCCA TIGGIGGCAG CGGGTACCNG TCCTTTNTIG TTCAACATAG GGTAGGIGG AGCCACGGGI CCAACICGCT TGAGGCTGGG CCCTGGGCGC TCCATTITNT MITCCAGGAG CAINIGGIIC TITGGCGCAA CCCACGCAGC CCTGAGGATT

SEQ ID NO:1266: (Length of Sequence = 322 Nucleotides)

CGGACAGATG TCACTCTCGC CCGAGAAGGG GGACACTGTG ATGGTGTTCT TAAGCTCATA GAGTGGCAGG TTGTCTGAAA
TGCCACCATC CACGTAGGGC ACCCCCTGGA GGGAGGGGG GATGAGCCCA CAGTACACGG GGATGAAACC NCTGCAGACA
TTGGCCTGGA TGAGCTCGTC CTTGGAGTTN AAGTGGGATA TAATGACATT NTCGCCGTCT GACACGCGGG TCAGGGAGAT
GCCCAGGCGC CCACTGGCAT GCTCATGGCT ATCAGCAGGC AGGACCTTNA GCAGGAAACT CGGGATGATC TTTTACCAGG
TT

SEO ID NO:1267: (Length of Sequence = 310 Nucleotides)

GTAACCCATC CCATAGGGIT GINCTATGIA TICTIGCCAG GIGGGGITGG AGCACCTTGI GAGCICAGCA GCCCAACATC GATAGIAAGG GAGICAGGGI TICTICATCI TCCCTAGAGI TAGAACTCAC TICTACAGCC ACIGIGICAG GGACCACTTT GAGCGCCCTT GGCACCTGCI GGCTGGAAAT CAATTTAGCI GTAATGGATC TGGCCCAGCT TITCCTCTCT TGGGTCATCT GCACTCATAG TGGTTGAAGC AAGATCTACC AGATGGGGAC ATTGAGATGG TCCCTTTCTC CTTCTCATTT

SEO ID NO:1268: (Length of Sequence = 338 Nucleotides)

GGGCTGCTCG TGAGGATGGG ACAGCATTGA CTTACTGGGG AGACTCCCTT GATGACAGCC TTACACGGTT ATTCATAAGG
AGGCAGGAAG AGGCGCTAAC AGTAAGCATG TTCTGGGTGG TCTTCGGGGT GCACATGTGC AGCAGCTGTA CCTGCTTGCT
TGTATGTTAC ATGTCTCATT AACATCTGAA ATCTCCACCC GGGAGTGTGT TTTTNACTAT TATAATGAGC AAAGGTTCAG
TCTGAGGACA GGTAAAATCA AAAATGTGCA CCCTCTTACG GGGGAAATTC CTTACTGGAG CTAGTTTGGC TTGAAGNGAA
CTGGACTACA GTGTGAAT

SEO ID NO:1269: (Length of Sequence = 363 Nucleotides)

CIGCTAGAGA GTATTTCAGG GTCTGCAGCA TGTTGTTAAG GCCATTAAGC ATATGTTAAG GCCATTAAGA GCAGTAATTA
TAAAAGGGCC CTGCTAAAAT AAATATCAAG TTCCCTTAAG AAACTTCAAA ATTATGAAAG TTTCAGGTCA TTATTTTGCT
ACAAATGANC TTAGCAGCTA AGNAAAATGT CTGCCTGCTT ATAAACTAAA TATGGTATAA TTATATATTN CTNTTATGTA
TTTCTAAAGC TACATTTTCA CCCTAACTCT ACTACAAAGT AGTTTCGGGA AACAAAGTAA AAGCAGGGGN AATCCAACTT
CAAATATAAT CAAATATAT

GATAAGIGAG ACTAATGGAA TCGITTCCCT CIAACTICAT AAAAACTITA AGGATTATCT TTCTTGAGIT CICTGIATTT
CIGITTIAGA AGAAAAGAAC AAAATTICAG AAACAAGATT ATAGIGCTIT INCIAAAGIA TAAATACGIG GGCCCTATAC
AAACTGGCAA ATTCATTAGT CITAAAGCAG ACATCCAAGC TATTGIGGGT GITTGGATGA CACCATTITC ACAGTAGGAA
ATCATTTCAT TCTGAGCGIG GGAATCGGCA TTGGITAACG CATGAGGTTT TATGIGGTAT AAACACCTGG GAAGTGAGAG
AAAAGNCAGC ACAGAAGCTC TGTGGGAGCT CITCTGAGCA TTG

SEO ID NO:1271: (Length of Sequence = 335 Nucleotides)

ATGCCTCTGG CTTGTTTGAC TGCAAAAGGT GATGTGCAGG GGTAGAGGTA GGGTACTAAT TTACAGTCAC CAAAATTAGT ACTGATATTA ATCAGTTTAG TTGGATTAAG ATGAACAATG TTTAATGCTT TAAGGAYCAT TTTTTTGCCCC AACAGGACTG TGCTATATTA AATGACACCG TGCCCAAAAAG CTCAAAAAATT ACATAGAAAG TAAAGTACTT CTTGAATACT AAAACAGTTA AGCATAAAAAG GTTGTGAATT GGTCCCAAAAG TGATATTAAC TTAAACATTT AATCCTACGN NCTATCTTAG CTGTACCCTC TAAAAAATGCT TAGGA

SEO ID NO:1272: (Length of Sequence = 323 Nucleotides)

GTTTTAGATA TTTTAAGATA TTTAACTGTC CCCTGTGGCT TTTAAGGAAA AAATAAGTAT AAATNCTTGA ATATTAAGAN TTTTAAATCA GCTAAATTCA GGGCCAAGAA CTATTTAAGA TGATTCANTG AGAAAGAAAA GGACCTAACC TGGAAAAAGA GTTTCAAAATA TGCCAGTACG TAGGGTATTT NIGGAAATAC ACAGTCTAAA ATTAAAAATT NNAACTNATC AATGGAATTT AAATCTATAG CACTTTAAGG CTGTGGAGCC CAACANTAGG GGNTACTTTG GGGGCACATG ATCTTTCAAA ACATAAATTA GGG

SEO ID NO:1273: (Length of Sequence = 368 Nucleotides)

GCAGCCTGGG CAACAAAGCG AAACCCTGAC TCAAAAAAAA AAAAAAAAA AAAGTCTCTT AATCACAACA GCAAAGCTCC
AAAAGTTCAA GCATCACAGG TAGCTAGTGG CTACTATATA GGNCAGCACA GACACAGAAC GTTTCCAACA TCACACACAG
TTCTANTGGG TAGCAATGAC CTATACTGCT GACCATGCTG NCCAACATGT NTGCAGCAGT CCCTCATCCC TCTGTNGTCC
CCTGTTACAA GCTTAGANCC CCCTCCNNAC GCTCCTCCCC CATAAACAGG GCAAGTNGGG CAGAAGGTGG AATCCTTTTC
AGGGGGCAAA T 331

GCAATGGGAT CTGGAGCCAA AGAAAAATAT ATCTGAGTTC TAGCTCCTCA CTAAGTAACT GTGTGATAAT GGGTATGTCA
CTCACCCTCT TTCAGCTTTG GGTCCTTTAT GTGTAAAAGG GAAAAACATA TGCCTACATC ATAAGGCAGA TGTGAACATC
AAATGTTATC AGTAACTGTC AATCTGTTTT ATTAATTGTA GAATGTCCAA AATATTAGTT TGTATGGACT TCAATGAGTA
TGTTTTGTGG AGTGGAGTGG GGGAAAGGGA TCATTGCTTA CCCTCTGCAC ATATCATGTT TCAGCCTAGT ACAAGGCAGC
CATGAGCACA AAGGGCTAAG CTACTTAAAT CAGNCCCCAA ACAACTTC

SEO ID NO:1275: (Length of Sequence = 319 Nucleotides)

AGATTACTCT TIGCAGAATT TIGGITAATT GIGAAGCIGA AATATCCIGA CICTACCICA AAGITAATGI TITAGGIAAC
TGAACAGGIA TICINCCCAT TACTAGTATT GAAGICAGAA TACAGAAACA AATAGTTACT GCCAGAAGCA GAATGGAAGA
GCCAAAAAAGI ACACAAAATG GACGCCATAA AINCIGAAAT AAAAGIGTAT GATGIGTICT GAGICACTGI AGAAGICATG
CATTIATTAT CAAGATAGAA AAGAGCAGAG AATGACGIGG GACATTGGIC CICGGAGGGC TICGTANGIG GITCGGICC

<u>SEO ID NO:1276:</u> (Length of Sequence = 324 Nucleotides)

CTGCATTGGG CAGGACAAAA CCTGCCAGAT TCAGAAGGTC ACGANTCATC TGGCCTTTAA TGCTGATATC CAGTGGAGAG CTGGAGTGGA GGCTTGGGGA AATATTGACT TCCAGGACCC AGGGCTTGAG GTTTTCNTCT AGCATGATGT CAAAACCAAA GAGTTCATGG CAGCTATAGG GCCGTCGCAC ATACATCTTG AGCAGGCTGG TCACATAGGG CTCTGACGAG ATGATAGTTT TGACAAACAAC ATCCTTTATC TTCTCCCAGA TGGCGTCGCT ATTGATNCC CTTCTGGGCT CAGGTAGTTC CACAAAAGCC TTCA

SEO ID NO:1277: (Length of Sequence = 388 Nucleotides)

AGCAAGGCGG TGGGGTAAGT NTGGACCTTT GTGTACCAGA GAGAACATCA TGGTGGCTTT CAAAGGGGTC TGGACTCAAG CTTTCTGGAA AGCAGTCACA GCGGAATTTC TGGCCATGCT TATTTTTNIN CTCCTCAGCC TGGGATCCAC CATCAACTGG GGTGGAACAG AAAAGCCTTT ACCGGTCGAC ATGGTTCTNA TCTCCCTTTG CTTTGGACTC AGCATTGCAA CCATGGTGCA GTGCTTTGGC CATATCAGCG GTGGCCACAT CAACCCTGCA GTGACTTTGG CCATGGTGTG CACCAGGAAG ATCAGCATCG CCAAGTCTGT TTTCTTACATC GCAGCCCAGT GCCTGGGGGC CATCATTTNG AGCAGGAATC CTCTATCT GGACITGIAC CCTGGGTGGT GAGAAGACCC TGATTGGTT TATTAGTGCA TTTCTGTAAG TNACTGGGAT AATCATGTTC
AGTTCAGCAT TTTATGTGAG TTTCTGAAAG CNCTTTAATC AACTCCATAG ACAAGATTAT AGTGTTGCAC AGCAATAGGC
ATGGGCCATG TCTGCACTGG AGGTAAGTTG CAAGGTACAC CCACGGGTGA TTTATCACTC TTACAAAGAT GATAACTAAT
GAAGACCGCA TCTAGAATGC TCTTACTGGA GATGGTTTAC AGAGCATTTT TAATCATCAT ACTTAGATTT ATATTAATAT
TTCTTTTCAA ACTAAATTAT TCCAAACTGT GCCC

SEO ID NO:1279: (Length of Sequence = 347 Nucleotides)

CCACTICAGI GCITCIGIGI CCCGAAAAGA TCTTITGACG CATAGGGCCI AACTGIAATA CACTIAAAGG ATAAGTCICC ACCCCAAGGI GAACATGGI CATGIGITAC ACGCACATTA GITCATTATC CATGIGIGAG GACCICCITI GIGAACAGIC ACAGCICCTC CIATAACCIG TTAAATATGI ATGITTGATC AACCCATTCA ACTTAAATNC TIGICITACC TCTCCITCCC TCAAAGTGCC TGGCIATACT TCCCAGCCTG CGGGATGGCC ACCTTGCAGG ATGGAACCCT TTGTAAGAAA TAAAGTCTCC TTTCCAAATG TACACATTGI ATGACTT

SEO ID NO:1280: (Length of Sequence = 344 Nucleotides)

ATCCTIAGCA TGCCTGINIT ACTGAGACCA TAAACTITIT INITITCCIT CTGCCTTCAC CCAGTGTGG TTAAGTCTTG
CTTGTTAAGC TCCCACACTT AAATGGCTGC TTGCAGAATT GCAAAGGGAC TAGGGAGAGA ACAAAAACAG ATATGCAGGT
GGTGGTTGTT AACCAGACAG GATTTCTAAG GAGGGTTCAG GCAGTCAAGT GGTTTTNTGT ATGTNTTTTA TGTTCATAGT
TTTGAGTTTT ACAATGTGG AAGCTTACTT TTGCTAGCAT TAGGTATAGT TTATTTTGAA AGAATGAGGC TCCTGAAAAT
AAACATGCCC AGTAAACTAT ATCT

SEO ID NO:1281: (Length of Sequence = 331 Nucleotides)

TGAGGAACAT AAAATGGCTT GGTAAAAGTA ATAAAATCAG TACAATCACT AACTTTCCTT TGTACATATT ATTTTGCAGT
ATAGATGAAT ATTACTAATC AGTTTGATTA TNCTCAGAGG GTGCTGCTCT TTAATGAAAA TGAAAATTAT AGCTAATGTT
TTTCCCTCAA ACTCTGCTTT CIGTAACCAA TCAGTGTTTT AATGTTTGTG TGTNCTTCAT AAAATTTAAA TACAATTCGN
TATTCTGTTT CCAATGTTAG TATGTATGTA AACATGNTAG TACAGCCATT TTTTTCATAT GTGGAGTAAA AATAAAATTA
GTATTTTTAA A

SEO ID NO:1282: (Length of Sequence = 310 Nucleotides)

CCATGICAAA TGTAGTTTAC AAAGGAAAG GACAAGTACC TTINTATAGA ATATACAGAC ACAGCATCAC ACCACAGGGC CCACGGGAG GTOGGGGAGA CGACACTTTT TCCCTGGGAA AGGCAGCTCT AATCCCAGGA ATGGTTCTCN GCAGAGGCTG GGTGGCCAGG AGCACTGTCC TCTAGCCCCC TAACTCAGCC TCTGCTTCAN CTCGGTTCCC ATTTCCTGCC TCTACCCCCC AACTCCTTAT AAAGAGCCCC ATGAGCTAAG ACTAAGGAGA GGTTCATNTC CCTTGGGGGG TGTGCCCCAT

SEQ ID NO:1283: (Length of Sequence = 323 Nucleotides)

ATGAGGATTA ATTATATCTG INTCACCCAC ACAGCTCCCC CATACCCATA ATCTTTATTT ATTINCTTCG TTTCTTCCTT
ATACCTTGTT TCAGGCATTA AACCATAACC TGITATTTAT NCTATCCTTT TCAAAACAGG TGIGGACCAT GCACAGATGA
CCTATGACGG GCAGCACTGG CACGCCACGG AAGCCTGCTT TNNTTGTGCC CAGTGTAAAG CCTCTTTNTT GGGATGTCCC
TTCCTTCCCA AACAGGGTCA GATTTACTGC TCAAAAACGT GCAGTCTTTG GGTGAAGACG TCCATGGCCT CTGAATTCTT
CCG

SEO ID NO:1284: (Length of Sequence = 283 Nucleotides)

TTTTTTCACA AGGTGAAAGA CCTTTATGGA CATGACAGAG AGGACCTGAG TTAAGAGGGA AAATACATCT NCATAGCTAG GTTCACAATC AGTTATGTTA GTCCCAAACC TACAAATTCA ACATGATCCC TATTAAAATC CTACCAATAT AGTTCAAAAG CTTGACAAGT TGATTGTNAC ATTTATATGA GAGANTAATT AAAAAAAAAA AAAATAGGGC CAGGTGCAGT GGCTCACGCC TGTTATCCCA GCACTTTAGG AGGCCAAGGC GGACAGATCA CTT

SEO ID NO:1285: (Length of Sequence = 341 Nucleotides)

CATTCINATG ATGTAGAGGC CAAAATGGTA TITNATAAAG AGGAAATTAC TTCTGANCCA CCCCAGCTGG AAACACTGGT AGTATCGGCA GCAGATGTGA TTACATCCGT TTTGGTATTA CACATCGTAT TTACAGCAGA CATGACTGAN CTGGGAACAC TGCCCTGTGA GACAGCCTGA AAGTTTTTTIN CAGATTTTINT GTGAACACTG TCTGAATTCA CATTTGGCAA AATGATTCTN CCAGTTTTCTC CGGCTTCTGC TAGTTTGAGG CAATCTGTT TATGTGCCCC AGCTGAAGAT CTTTCACTAA CTCGATCTTT AAGAGCTAAC TGCATTGCTG G

SEO ID NO:1286: (Length of Sequence = 354 Nucleotides)

GCCCTATTIG TACAAAGIGI GCATGINAGC GIGCGIGIGI GINTIGCATI TITICCCCCIT TAGGIGGITC AAATITIGGAA
TITIGIGAAGG CAGAGCIGAT AATTAGAGAC AATAAAAATC IGCAGAGIAG AIGGITCCAC AAACAAGACT AIGAAAGAGG
GGATAAAAGA AGAGGICAAG AAAGACTCAA GAACAGIATA TAGAAATAAT TCAATTACAT TAIGIGIATI TITAAAGAAAA
CATGITCAAA CIGCATGAGA CAGAAAATAG CACICNGITA TCCICCIAGA CITCINAAAG TITIGAGITT GICIGCAATC
TCTICCATT AATCGNCIIT TGCCATCTIC AGAA

SEO ID NO:1287: (Length of Sequence = 354 Nucleotides)

CTCTCTCACC CGGTGGCCTA TAGCCCCCAA CGTGGTCAGC AGCTGCCTCA GCCATCCCAG CAGCCTGGTT TACAGCCCAT
GATGCCTAAC CAGCAGCAGG CGGCTTACCA AGGCATGATT GGGGTCCAGC AGCCACAGAA CCAGGGCCTG CTCAGCAGCC
AGAGGAGCLAG CATGGGGGC CAGATGCAAG GCCTGGTGGT TCAGTACACT CCACTGCCTT CTTACCAAGT TCCAGTGGGT
AGTGACTCAG AAAATGTGGT CCAGCCGCCT TTCCAGCAAC CCACTGTGT CCCTGTGAGC CAGTNTGTGC AAGGAGGCCT
NCCAGCAGCG GGGGTACCA GTGTACTATA GCAT

SEO ID NO:1288: (Length of Sequence = 231 Nucleotides)

TITACITAAT TGGTATAGAT TGAGGNTCAT GCATCANCAA GCAGTTTTGA AATTNTCCCC AAGTGATTCT NACCTGCAGC CTGGGTAAGA AGTCGCAGGG CTCCTGGATA GTCATTAAGT GAACTGTGGT AAGCACTGAT GTAGCAGGAT TACCTGCCCT ACTAGGTGCC GGAACTGCAT TINCTTGCTC ACAAGTAATT TTTTTAAATG TATGCTCGCA TCCCTGCCTT G

SEO ID NO:1289: (Length of Sequence = 329 Nucleotides)

GGACACTGTG AGGGGAAAGG ACAATTTTAA AATTCCTTTT CAAGGAAAAA AAAGGTCTTT ATGCTTTGCC ATGAGGCCAC
ATTCAGCTGC TATTTAANCT TAATATCTTG AACCTAAAGA ATGCTGACTT TNCCTACATT TCCAGAGTTA GGCAGTATTC
TACACTTAAA GACTACTACT ATTTTNATAA AAGGTAATCT ATTCAAATTT CTTCACAGAT TTCCCTTGCT GGGGATCAGT
TAGTAAAGAA GGAGGAATTC CTCTTACCCA AGAGGAATTG CATTGCTTTA ATTTAGCAAT GTGAGGTAAG GCCTGCCNAG
TGCCCAGGG

SEO ID NO:1290: (Length of Sequence = 297 Nucleotides)

GGAGGCACAT GTGCAGCTTT GTTTCATGGG TAAATTGCAT GTTTCTGGGG CTAATGGGT TTCTTTTACA GAAAAAAGTA
TCAGAAATAA TCGGTTAACT TTNCTCACAT GGTCTTAACT CTTCTTCAGG AAATATCTAA CTTGTAAGTG CAATCCTTCT

TGTATAGCTG CCAGACCAGA CCCAGATAGA CCATAATAAA ATAAAATACA CAGTCAGTTT TTAATGCAAG CCAGAATGAC TCINCTGTAT CTTTAGCCTT TCCAGGGGGA TACAGTGAAC TCAGATATCC CTGCTTA

SEO ID NO:1291: (Length of Sequence = 317 Nucleotides)

CTATAATCCC AGAACTITGG GAGGCCGAGG TGGGCGGTTC ACCTGAGGTC AGGAGTTGAG ACCAGTCTGG CCAACATGAT
GGAACCCCAT CTNTACAAAA ATAAAAAGCA AGAATCACCT GAGCCCAGGG AGGTCGAGGC TGGCTAAAAA TAGATCTGGG
GGTAGTGGTT AATNGGGCCT TGTGAATNAT TCAGCATAAG GAACTGTCCA ATATTTTTTT AAGCTGTCAG AAAATCC

SEO ID NO:1292: (Length of Sequence = 293 Nucleotides)

GAAGATGGAA ATAGACCACC ATACAAAACA AAAAAGACAG AAGAGAATAT TAGCACTCTG TTGCAAAGGA GAATAGGTAT
GCTCAACTGG TAAGTAGAAT GCAAATATTC CAATATCTGA AAAAAATCCC AAATCCAAAA TACTTCTGGT TCCATGCATT
TTINCTAAGG GATACTCAAC AGGTATTTTA AAAGATCAAA ATACAGATCA GAGAATATGG ATACTTGAAG ATTATGAGCA
AACGAGGATT AAGGNAAACA TGTTGGAGGA CTTTTTAAAA ATGTGTTAAA GGG

SEO ID NO:1293: (Length of Sequence = 310 Nucleotides)

TCCCAGAAAC ATTACGGITT GATATCAAGT TCCTATTITA AGAGTCACCC ATTIGCCCAC CATAAGINCC TGGAGAAGGT AGGGTATTAC AGGACTAACC TTCCAGIGGC TGATTCTGGT GGITTCCACA TTCAGGITTC TCTGATTITN ACAAGCTITT TCCCCATAAAG ACTGCATTIN CITTAAAAGC TTCTCCTGCA AAANAGCCAT AAATTGAAGC ACCAGTGAAG ACAATAAAGT AACATACAGA CCGTTTCATT GGGAGGGGGC CCNGAATGNG AGACAAATAA GTCCCTAGTA AATGGCATTT

SEO ID NO:1294: (Length of Sequence = 275 Nucleotides)

GAATGACGAT GTCAGGGGCA TCAGGAAAGG TAAGGGCCGG GAAACCGGGC CCTTGGGAGAA CCCTGCCCAG GGGAGGCCCA
GCCTACTCAC AGGNTCCGAC ACTCCAGGCA GAGCAGAGGG CAGGAGAGGC CCAAAGAGCT AGGTCAAGCA GCTGGCTCCC
CTGGGGTTAA ATACATGGGT TTTTGTTTTA CTGCTGTGCT TGATATACAT GAAGTAATGA ATACAAGCA ATTCATTTTT
CCTGCATCTT TACTTTTACA TTTGTNCTTA GGTTGCCTAA AACATTINAA ATACAATAAA ATGAGTGTAG CAAAAATTAT
TGAAGCT 327

CAACCTCIGC CICCCGAGIT CAAGCGATIC TCCTGCCTCC CGAGIAGGIG GGATTACAGG CATGATCCAT CACGCCCAAC
TAATITITIA TITITAGIAG AGATGGGGIT TCTCCGIGIT GGICAGGCIG GICTCGAGCT CICGACCTCA GGIGATTCAC
CCACCTCGGC CICCCAAAGT MITGGGATTA CAGGIGIGAG CCACCGCGC AGGCTACTGG TCTCAATTCT TITGGATACC
CAGAAGCAGA AATGCIGGGA TCACATGGIA GICTC

SEO ID NO:1296: (Length of Sequence = 247 Nucleotides)

GGAAGGAACA ATTGATAAGA ACCGGGGACA TCAGGGAGAG AGAGTATTTG AGCTGGGCTT GATTCCATCG GGTAGTATCT GGAAAAAAAA AAAAAATCC CAGATGAAAG AATGTACAAA GACATGAGCA TGCAGGGCAC ACTTTGGAAA ATGGGNGAAG TCTGACAGGC CTGGGAGAAT GAAGACAAGT TAGCACCAGN TTNAGAAGGC CTTGATTACA NGGCCAAAAC TTTTGGATTT TACACTA

SEO ID NO:1297: (Length of Sequence = 246 Nucleotides)

GACTICITAC AATGCAGCAG CAAGAGAAAA TNAGGAAGAA GCAAAAGCAG AAACCCCCAG TAAACCCATC AGACTICGTG
AGACTIATIC ACTATCACTA GAATAGCATG GGAAAGACCA GCCCCCATAG GTCCACTACC TCTCCCTGGG TCCCTCCCAA
AACATGTGGG AATTATGGGA GATACAATTC AAGTTAAGAT TTGAATGGGG ACACAGTCAA ACCATATCAT TCTGCCCTTG
GCCCCT

SEO ID NO:1298: (Length of Sequence = 263 Nucleotides)

CATTGCACTC CAGCCIGGGC AACAAGAGCA AAACTCCATA TCAAAAAAAA AAAAAAAAA GAATTGCTGA CCTTTATGTG
TTTCTGTTTA AGTTCACAAC AGTCATAATT CTGTAAAATA CAAGGCAAAA CTGTAGTTTC TGATACTAGT AATATATCTA
AMTCAGTAAG TAAAAAGGAT GTGTAAAATC TTAATAGGGG AAATAATTAT TGTATGANCA AGCAATTTCA AAATCAAAAAG
NCACGTTTCA GTATATATTA TAG

SEO ID NO:1299: (Length of Sequence = 272 Nucleotides)

ATCINATIGI TGIGIAGITI ATGGCAGIGG TCTCCAGACT TITTGGCACT AGGGACCAGI TTAATGGGAG ATAATTITCC CATGGACGAG GGGATGGGGA GGAGGCAGGG GTGGITTCTG GATGAAACIN TTCCACCTCA GAAGATCATC AGGCATTAGT TTCTCCATAAG GAGGCAAAAC CTAGATCCCT TGCATGCACA GITCACAATG GCACTCGTCG CATATNCCGT CGACAACCCT TTTTTGAGGT TCCATGCTTC CCATTTGGCT TT

SEO ID NO:1300: (Length of Sequence = 277 Nucleotides)

ACCACTGCAC TCCAGCCTGG GTGACAAGAG TGAAACTCCA TCTTAAAAAA AATGIGTAAA ATGAAGATTA TCATACTACC TACATCATAG AATTGTTTTT AGIGTAAAAT GTGTGTGTT ACATTTATGT AATAGTTAAC ATTTAAAGAG CACCTACTTT GTGTAAACAT ACTTTGTATG AGATACTGTT CAAATATATA TNCTAATATA TGCAACATAT TATATATGTN AGAATAGGGT CTTATATATC TTAGGAAGTT AGATCTTATA TGTTTGA

SEO ID NO:1301: (Length of Sequence = 304 Nucleotides)

GGITGCGGGT TATGTAAATC CCAAACITAT GAACAGGAAA TGTGTACAGT GCATGATAGG TTAAATITTN CTTTATTGTT GTCCAACGCA GGTCCTTTGG AGAGAAAAAA AGATCACAGT GCTGACCAGG TAACTCAATA GGTTAAGTCA AGGTAACCAT TGAAAGATAA TAGGATTAGG GAGGTGTTA TTTTATGGCA TCTTCTCTCA TGGAGTTCTT AGCACTTCGG ACAATTTGTC TNTTCCCCAC TTTGTACAGC TGTTATGTGT CATTCACCAG CCGGCTGTAT TTAACTTGCC TACT

SEQ ID NO:1302: (Length of Sequence = 335 Nucleotides)

AGITTATIGO CATACAGAAA ACATITTATA AAATAATAIG GIAGACITCI ACITCAACAT ATICACGIAA AAACATCACA GIGCAAGAAA GIGATCACAA TIAAGCAIGA AGACATCAAA AGCCAGCCAG TATITIAACT ACAGAGCAGA ATATICIIGO TGICCCIICO TAGAAAAATGI TGGCACATIC ATIAACIGCI CAGGITACAA AAATCACIIC GIGICCACII CCIGICCIICO AATATATIIN CATAACIACA CIGIGITACA TIAATGCIGG TGGACAAATT AGCICCIATA AAATCIAAAA ACCITITCAG GIGGGCACAA TGGIT

SEQ ID NO:1303: (Length of Sequence = 316 Nucleotides)

TGGAGCTGTA TATGGTCCGG AGITATATGC AGCATCCAGC TITCAAGCAG ATGINICCCT AGGCAATGAT GCAGCAGTGC CCCTATCAGG AAGAGGGGGT ATCAACACTT ACATTCCTIT AATCATTCCT GGCTTCCCTT ACCCTACTGC AGCCACCACG GCAGCGCCTT TCAGAGGAGC CCATTINAGG GGCAGAGGGC GGACAGTATA TGGTGCAGTY CGAGCGGTAC CTYCAACACC CATCCCCGCC TATCCAGGTG TGGTTTACCA GGGACGGATT TTACGGTTGN TGACCTCTAT ATAGATTCTG CAAACT

SEO ID NO:1304: (Length of Sequence = 211 Nucleotides)

TATTITING TICTICTCC CCIACATATA TICTAAACCT TCIAAAGITT TIINATITIT TTAAGGATCA CITTATCATA
AAATAAAATA TCCITTTCAT ATAATAAATT ACCTAATAAA AAGTCTTTT TITTCATATT AGCCCAGGIN CITTGCTACA
TITTATATGGT AATAAACGCC TITTATTAAAA TAGANTATTA AATTATAAAG A

SEO ID NO:1305: (Length of Sequence = 316 Nucleotides)

GAAATGATTC AGGGAAAAAA ATTTATAGIA CGITTTCAAC TITTITITT TITCITTGAA ATGGAGIATG GICATAAAAA GGACACIAAA TAACCIGATI AAGCIAGAGI ATAGACCAAA TIGCCACITA CITTGAATIG TITTIACCAA AGGIATCACT TIGAATAAAG ATAACITTCA TIAGACATCI ATCITTATGI GITCCIGCCA TCATTTCAGI GAGATCAGAG GAAAGITAAA TIAGGAACAA TGAAAAAGCT TAAGAAATGA ACAATCATCA TGCITTIGIG TATGCITAAA GIGAGIACAT GIAAAA

SEO ID NO:1306: (Length of Sequence = 310 Nucleotides)

GGGGATTTTT GAAGGCTTCG CTGTGGATGG CCGAGAACCT GCTCGGGGTG TAGGTCTGTG TGTCTGGGGG ACAGTTTCCA
CATCTGAGCA CACGGACTGG ATTTCTGAAA TGTCAAAGTC TGATGCATCA CTGCCTCGGC GGCTGCTGGC CCINCTGCCA
GCTTTGCTTC CAGCTCGACT TCCTGGTCGG CTGGGAGTCT TCTTGGAATC AGCAAACTGT GTTCGGACTC TGGCAGNTGC
AGTTGTTATC AAGCCACTGT CCTCCCCANA GTGGAAGCCT TTCCCTGATA AAAATCCTGG AAGTCGAAGC

SEO ID NO:1307: (Length of Sequence = 302 Nucleotides)

TAATAAATAG TATATGTAGT GAAGAAAAAG TTATAACAAG TATACATTAC ATTTAACACA CCTAGCACAT AGGACACCCT CAACAAACAG CTACAGCTGC TGTAAATCAT GTGTATATAA TATAACATGC AAGCATATCT TCATGTATTG ATTAATTACT ACTTCTTGGA AAAGGATCTG AGGAACATAT TTAATATATT TNATATGCCT GCTCATATGT NCATTTAGTG CTTATCAATT ATATTTAGTG CTTTTCTATT AGCTTCATCC ATTTGATTAA GATAGCAACT TGTATTATTT AA

SEO ID NO:1308: (Length of Sequence = 285 Nucleotides)

COCCEGCCAA CGIGGICITC CICIACAIGC TCIGCAGGGA TGTTATCICC TCCGAGGIGG GCTCGGNTCA CGAGCTCCAG GCCGICCIGC TGACAIGCCI GIACCINICC TACICCIACA TGGGCAACGA GATCTCCIAC CCGCTCAAGC CCTTCCIGGI GGAGAGCIGC AAGGAGGCCT TTINGGACCN TIGCCTCTCT GTCATCAACC TCATGAGCTC AAAGATGCTG CAGATAAATG CCGACCCACA CIACTTCACA CAGGICITCT CCGACCTGAA GAACG

SEO ID NO:1309: (Length of Sequence = 319 Nucleotides)

TITICCAATTA TIATTITGCC AATATOCICA ACTCITITGC CCACTITNAT CITICCATICA ACCCICCIG CAAAATCCIG ATCIAAAAGC AACCCAAGIA TITIGCCICTI CAACCICCCA GCIGCTEAGI GGITITGGGA ATTACACAAC CACTAAGCIT GGIGCAGAIG CACTATGGCC TCAATAGAGI CCCCAGIGC TGCCCACTIT CICCITCCAT ATTICICCAC AGCAGCIGGI CAAAAATACAT TINICCCCAA ATGICTIACA CAACCCCCIT CICCITTATC ATCCTTANCI CACCCCCACC CCAGITCIT

SEO ID NO:1310: (Length of Sequence = 356 Nucleotides)

TGAAGTTTTG CTCTTGTCGC CCAGTCTGGA GGGCAATGTG CGATTTCAGC TCACTGCAAC CTCTGCCTCC CGGGTTCCAG
CGATTCTCCT GCCTCAGTAT CCCAAGTAGC TGGGATAATA GGCACTTGCA ACCATGCCCA GCTAATTTTT GTAGTTTTAG
CAGAGACGGG GTTACACCGT GTTGGTCAGG CTGGTCTTGA ATTCCTGACC TCGTGATCTG CCCGCCTCGG CCTCCCAAAA
TGCTGGGATC ACAGGCATGA GCCACCGCAC CTGGCCCTAT ATCCTGCTTC CTATCTCGTG GGTCATCGTT TATCCCTTTT
ATTTATTTCA ACCTGCAGTT GTTTGCAGAA CATCTG

SEO ID NO:1311: (Length of Sequence = 331 Nucleotides)

AGCTCAGATT CATGTCTTGA GCCAAACAAG TGAATGTATC TNAGAAGACT CAGTACCACA TGGTACTGGG AGATCTTACT
CACTTCAGCT GGCGTTGCTC ATTAGTGAAT GTATGACAGC AGGATGTGAG GGGATGCCCA GGAGTCAGTG TTAGCATTGT
CATCTGAGAT CACTGCTATT AATATCATCC ATTAATTTAT TAGTGAGCTT CACTATATGC AGACTGGGAG ATTAGGAGAA
AATCTGTCAC ATTCTCTCTA GCTAATCAGA TCAGCTACCA ATTAATGAGA TTCTGAATGA AATATCAATA TGTGTTTTTC
TAATTTGGAC C

SEO ID NO:1312: (Length of Sequence = 347 Nucleotides)

TTTTTTCCTT TATAAATTAC CCAGCTTCAG ATTTTTTNAT AGCAATGCAA AAATGGCCTA ATACACTTCA GAACCTGGAA
GATTAGCAGT GAGAAATAAA ATCAGTTAAG TTGATGACTT CTAGTATTTC ACTACATGGT TGTTTTGCCA AAATGAAGGC
AATTATCAGTG TCTTCACACT TAAAAAGTAG TATATTGANC TTTGAGGTGA AAGAGCTGGG GTTTAAATTT GINCTTTACC
AATTATTGAG ATAAGTGTCC TTGAGCAAGT TACTTGCTTT CNCTGATCTT TAGTTTTCTT ATTTGTGAAA TTGGAAATGG
TGGTGTTTCA GAGGGGGGTT GTATATA

SEO ID NO:1313: (Length of Sequence = 336 Nucleotides)

GAATTCCTCT ATCAAAGTGT TCATAAAACC TGGAGCTGCA GCTGGCCCCC ATTAGGTAGT TTCTTGGTGA ACGTTTTCCA
AGGAAAACTT TTTTTTAACA ACTTCATAAA GCCAAGCACA AAAGGACATT GCAATGACTG GCTGAAAGAC ATGGGACTTT
TTGTCTTCGA CGACTAAAAC GTTAAATGGG GGCTTACTTT GTGCATTTAT GGAAGAAAAC TTGGAAGGCA TTAAAGGCTA
CATTTTGAGC CTTGCATGAT TTCATTCATT TATGCATGAA TTCATTTGTT CAACATTTAT TTAGTACCCA CTATATGCCA
GGCACTGTGC CAAATG

SEO ID NO:1314: (Length of Sequence = 391 Nucleotides)

COGGITTAGA COTCAGTOGG OGCTGTGAGG GCACTGTCCG COCACCTGCT OGGCTGGCTG AGCTAGGTCA GTGGAGAGAA
GCTGGGGCCA CTCACACAGC ACAGCAGGCC ACAGTCTACA GAGTACGCCA GGTAGAGCGG TTAGAGTGGC AGCCGCTGGA
GAAAGGGTTA TAGAAACACA TOCCTGACTC TTTGGTTATG TCCCACGTCC TCTGTGTCTC CTTCCCCTTC CCTACTCTCC
TTCCTTTCTG CCTCCCTGCC TCCCTTGGAA GTCCCTGTTG TCAGTGCATT TNAGTGCATT GACGTGTCCT AAACACTGAT
CTNCACACAC CTTCTTTAT CTTCCACCTG ATAGGCAGGC CCCAGANCCC CTTTTTTCCT AGCTTTGTTC T

SEO ID NO:1315: (Length of Sequence = 374 Nucleotides)

GAATTCCCTG GAACACTGGT GTTTACAGAG AGAGATACTT TGTGGAATGG AGCTTACATG ATGAATGAAA AAGAGACCGT
TAAAAAGTAC TAGCCGTTGT TTACAAATAA CTACCAGGTA AACAAAGAAA TCACTTTCTT TCCCCTTTCT AAGGATAAGG
GAGAATAAAA TAATCACCAA GAGGCATGGA GTTTGAAAAG TATATAACAG ATTTCTTTAT TATTATTTAC AATCAAGTTC
TGTTGGNCAA CATAATGAAA TAAATAAAAG ATGTGCCCTG GCCTGTGAAT TTCAACTCTC CTTGACTTAA GTTCTCTGAA
GGGCAAATTG GAAAGCGGTG ATCAGGCAGG GAAGAGAGGG CCAG

SEO ID NO:1316: (Length of Sequence = 353 Nucleotides)

CTTGITTACA GGITTIGAAA GGITTGINAG ATTAGTATIT ACTITTAATT TITTGAGTAA TAGAATGCGI TTAGGITCTA
AATTACTATG GAAATGGCAT AGTGAGGATT CINCACAGAT ATTAGAGACC TICAACAACA TAGTGAAAAT AGATTTGTCC
TTTCTTGIAA ATAGCTGAAC TATGAAAATT TGANCTGICA CTGGAGGGGG CATTTGCNCT GAAGTAAAAA
TAACTTINCT CITTAGTAAG AAAAAGCTAT ATTTINCAAT ACTGCCTGCC ACAGCAAACA AACAAAGTCT TGTTTGTGTT
TTAATATTGG CAAAGGAAAA ATTCTCTATA TAA

321

SEO ID NO:1317: (Length of Sequence = 316 Nucleotides)

GAATTCCGAT TATAAGCATC AATATGCATA AAATGCTTAG AGATGGACCT GATATATAGC AAACACTCAG TAAATGTTAA CTATTATTAT NACAGCACAG CAATTTATTT AAGATTACTG AGTGTTCAAA TGAAAAAAAA GACATATTAA CTTATATAGT GCCATTTCTG ACATAAGAAA TACACAAATA GAGGTAGTTT CTGAAACAAA GATCAAAAAA ATCTATTGTA TGGTGTCTTG TATCAATGTG GCTAAAATTT TCGAGCTAAG TTTTATNAAA GACAGATCAT ATTTCANGTA GGTGATTTTT GTATTG

SEO ID NO:1318: (Length of Sequence = 300 Nucleotides)

GIGGGACTAC AGGIGCACGC TATCATACCC AACTAATTIT TGTATTITTA GIAGACATGI GITTCCCCAT CITGGCAGGG CIGGICIGAA ACTCCIGACC TGAGGIGATC CACCIGCCIT GGCCTCGCAA AGTGCIGGGA TTACAGGIGT GAGCCAACAA GCCTGGCCCA TTTATTIACT TITTAATTIT CATTITTCIT CATCATGIAG AATGGACAAT TTCAGGAAAC TGATAGAAAA TACIGICIAA CATCAAATTI TCAAAAAAGT TTCICTGTAA CAGATAAGGC AGTCAATTIC

SEO ID NO:1319: (Length of Sequence = 306 Nucleotides)

CAATAAGCTT TAAAAAGTTA GTGCCACATG ACCAGCATCG ACTGGCCTCA GACATCTGCA AGCACTCACC CAGGCCACAG
GGTCAGGTAG AGGGCTCCTG GGCCCACTGT AGCCCCTGCT GGGTCAGTGT AGCTGGAAGG CTACGGGNCC TTAGTGGGGA
GCCCACAGCCT TTCCCACTAG GGGCCTCTC ACTCTGACAT CTCCCTGTGG TGTTCGGACC AAGGGTGGGG AGGGAGACAC
GCTGGCCCTA AAGGGAGGTG GTAATNAGTG AAGATCTCCA GGGCCAGNCC ACAGGGCTCC GTCCAT

SEO ID NO:1320: (Length of Sequence = 373 Nucleotides)

GGICTIGATC TCCTGACCTC GTGATCCACC CGCCTGGGCC TCCCAAAGTG CTGGGATTAC AGGCGTGAGC ACCGTGCCTG
GCCGAGATAA TTATTTTINA GTGACGATTT AGCAACCTGA AAACCTTGGG TCTTTGGGAT ATGACCTCAG TATCAACACA
GAATATTTGA ATGCTGGTTA ATATATTINT TTTAAACTGT GATAGAATTG AAATCTTGTA GCCACATTTT GAAAGTTTAT
TCTTCATTAA CTAGTCTTTT CTCACCTGAT TTTCTACAAG AGAGAATTTT CCAAAAGGTT AGTTGTCGTT ACATTAAGAA
CTTGGGGTTT GNTTGACATG AAATGTTTCT ACACCAGCAG GTCTCAGATG AAT

SEO ID NO:1321: (Length of Sequence = 366 Nucleotides)

GITTGGCTAA TCATCCTATG ATTITCCTAT AGCITGAAAA CITTTTATAT CITAAATTIT TINATAATTI TGAAGTATTA
TTGTTTGGGC TITGTATATC CAGTGTATTT TCAATTAAAT TCCCCTAACT AAAGTAATTC AAAAGGAATA AAAGTGATAAT
GIGGGCTGGG CGTGGCGGC CATGCCTGTA ATCCCAGCAC TTTGGGAGGC CCAGGCGGGC AGATCACCTG AGGGCAGGAG
TTGGAGACCA GCCTGGCCAA CATGGTGAAA CCCTGTCTCT ACTAAAANTA CAAAATTAGC CGGGTGTGGT GGCACATGCC
TATAATCCCCA GCTATTTGGG AGGCTGAGTC AGGGAGAATC TCTTGA

SEO ID NO:1322: (Length of Sequence = 362 Nucleotides)

AGGAGGGIA AAACAATCC CCCICCAATG CTITGIAGAA GGGGATTAGA ATCACTGIGG AATTCGGIAT TGGCTAATAA
AGTATAAACG CTAAAGATCA ATGCCIGAGT GCACAGITGT CCTTCAAGCC ATTGIACTIC TGCTTTCCAA GANTAGANGA
CTACITTITIA ACCAAGANIT AAAAATAANC TCATAATTTA AACACCICTT TCATGCCAAA TGGAAATCTT AGTGITGAAT
AATCAGGCTC ACCTGAATAC AAAGTTGICC TGAAAATGCT GACAATCACA AAAAAGGTTC TAGAAGCTTT TTCAAAAAAC
AAGTTCAGAT GGTTCCCACT GAGTTACTAT TTGAGGTTAA AG

SEO ID NO:1323: (Length of Sequence = 244 Nucleotides)

CGACCTCAGT GTAAATCACA AAACGGGAAG AGCTGATATT GGCAAAATAA TTACATGGCT CATTTCCTTG CATGTCAAAA
TAGGATTTGA TTGGTTGTAA AAGATGACAA ATACCTTINC GGTTTCAATG TTCTTAAGTG GGAAGTCACT TATTACAGAC
CINATTGGGA GTAAACAAAG CTGTTAGACC TTTCATTATC AGTCCNNTTA ATCCCTTCAA TAATCCCCCT AAATCAGTGA
GGCG

SEO ID NO:1324: (Length of Sequence = 279 Nucleotides)

GATCCATGCC ACAGTGACCT CTGTNACCCT GCACAGCACA GAGGGGAAAG CCCTGTACCA GGTGGCGTAT GAGAATGAGG
TAGGCAACAG CTCTGACTTC TATGACATCG TGGTCATCGC CACCCCCTG CACCTGGACA ACAGCAGCAG CAACTTAACC
TTTGCAGGCT TCCACCCGCC CATTGATGAC GTGCAGGGCT CTTTCCAGCC CACCGTCGTC TCCTTGGTCC ACGGNTACCT
CAANTCGTCC TAATTCGGTT TCCCAGACCC TAAGCTTTT

SEO ID NO:1325: (Length of Sequence = 338 Nucleotides)

TCAGITATIT GIGIGIGIG GIGIGIGIG GIGIGINCAT CIGCAAACCC TGCACITCAT TATCCAAAAA TIATITGATA
TITITATAATC AGAGAAAATG CIATITITAA ACCCTACCAC TGCIGACCAA ACAACAATCA CAACAGCATA ACACTAAATA
CIGITCAACA AATCTATITT AGIGIAGIAA TIAAATAATT CCTAAAATTA TAGACATCCC TAATATTCTT TCCNITIAGIG
GITCCTCAGA GIGCAATCIG TGGAGCAACT ACCTTGAAGA AATTTGGGGG AATGAGACCN TGGGAACCCT AAATGTTTAG
NATGGIGCTC TNGGGGAC

SEO ID NO:1326: (Length of Sequence = 393 Nucleotides)

AACTITIGAG GGGACACCAT CACTCAAACC ATAGCTGTAA ATCTATTCCT TGAGTCCAGA TCACAAATTA CCAAATGAAC ACGTTCTCCA TITTTAGTAC TTTTTTACCT GTAACCCTCT GTCTACCTAA GATGAATATT TATTCATTGA ATGAATCATT TAATTTTGGT GCCCCAAAAT TCTCAGTGAA ACAATTTCTG GATACCTCTC CATCACTAAG ATAATCACTA TAGCAGTGTC ATATTCTTCA ACTTAGAACA AATCTAAAAG CTCCATTTAT CCCTACTAGA AGTGTTCTGT TGTCTTTTTC ACTCTCAAAA TATCCTCCAT GCGCAAACCA AACACTAANG GGNACCACCA TATCTTGCTC AATGGAGGCN AAATCACTTT TTA

SEQ ID NO:1327: (Length of Sequence = 381 Nucleotides)

CTITGGAGAA TTAATTCAGC AGTIGGIAAA ATCATTCTAT AATAATGGGT ACCATTCTGC TCTGTCCCAC ATTITTATGA
AGTCTCTITA AATTTAAAAA GGCAATGIGC TTTGTGGTTC TTGAGCAACT TAAATACGIT GCTCTGAATA GTTATTGTGA
TGAGGTAATT TGTAACAACT TTTAGGATCA ATGCTAATTT NCTTAAATGT TTCTGTAGTT TCCCCTTTAT TATAAAGTAT
ATTAGGCTGG ACTCTTGGCT GTAAGTGGCA GAAAACTCAA CTCAGATTAG TTAAGAAACA AAAGGGTGTT GGTGACAGTG
GTGGCTTTCA GACTATTGCT GCAGGCCCAC CTGCCATCCT CTTACACCCT CAACATACCC T

SEQ ID NO:1328: (Length of Sequence = 289 Nucleotides)

AGAAGAAAAT TCTTAAGCAG AGIACTTAAG TACAAAATTG AGTGACTGAA AGATGCTTAA TCTAGGGAAA TTAAATGAGA
AAAATACATG GIGIGTGINI TGGAGGGGGA GCTGGAATTG GAATGGGCTG GAGTGATGAA AAAAAGCCAA CAGATATAGT
CTTCTGTTTT GTAATATAGG CTCAATACTA AATTATGTAG GACTAGATAA TCTAGGTCCT AATGTCTCCT TTTTGCTGGC
AACCTGGGGG CCAATTACAC TAGAGGGTTG GTAGAAAAAA GAGGAATAT

SEO ID NO:1329: (Length of Sequence = 364 Nucleotides)

TIGIATATIT GGGATVETCA ATAATCTAGG CCACGIGGAA GATAACAGGC TATTITIGGAT ATMINICIPAT TGCAATCTIT ATATTICIGT GTAAATGCCT ATACAAATGT TIGCTIGGTG ACATATGGAA AACITAAGGN CITITATGAA AAGGCGACAA TGGGGACCTC CAAAGCGCCA AAGITTCIGC TAGGCATAGT GTTATTTTTA GATTACATTA AAATGGCTAT TTAGACCCAT CTAGCTGAGA CTATTCCAAA ACAAACTTT TATCANATTG TNATCATAAT CAACTTTCTA CAGGCTAATG ACTTTATAGN TTTACTNCTA GTGTATATCT ACTAGCACAA TTGGACCCAG TTCC

SEO ID NO:1330: (Length of Sequence = 221 Nucleotides)

CAATATITAA ACAAAATGCA AAACTGAACG TTACCTCAAA ATGAAACAGT GTGTGTACTG GCTGTTAGAA GTTGATGGCG
GTCTACTGTT TGATATTCAC TGCCATCTTC CTCTGCCCCA CTCTACCTCA ACTCGGGACC GCCTCACCTA ATGGTGGGCT
TTGCCGCTTT ATGCCNTGTA GAGNAGACAC TGGGTAACCA CAGCAAATCA ACACGGGNTC C

SEO ID NO:1331: (Length of Sequence = 279 Nucleotides)

AATAGAGATA ATGGTCAACT CTTGAGAAGA ACCAAATGCT GGTGCCATCT TGGAAGTGCT ACATCACCTC CTCCTCTTAC
TTCCTTGAAC AGCAATATTT CTGGATTTCT TCTGCAAGCC CCAGGCAGTG CAGGATGCGT TINTTTTCAG CAGCCAGTTC
CTTCTCAGAG AACTGGCCCA AGAGTTTCTG GACAAATATA TTTTGATCTT TCAGAAATAT GTTCTNATTC ACTCCTACAT
TTGGCACATT NTCCAAGGGC CCAGACTTGA AATTGGAGG

SEQ ID NO:1332: (Length of Sequence = 290 Nucleotides)

GEACGAGGAG ATGTCTITGG TGGACTTGGG AAAGAGGTTG CTAGAAGCAG CAAGAAAAGG CCAAGATGAT GAAGTGAGAA CGITGATGGC AAATGGCGCC CCATTCACCA CAGACTGGCT TGGAACATCA CCCCTCCACC TTGCAGCTCA ATATGGTCAT TATTCCACAG CAGAAGTACT CCTTCGAGCA GGTGTTAGCA GGGATGCCCG GACTAAAGTA GACAGGACCC CCTTGCACAT GGCTGCAGCC GATGGACATG CGCACATCGT GGGAACTGCT TNTTCGGAAT

SEO ID NO:1333: (Length of Sequence = 201 Nucleotides)

CGCCCAGCIA ATTITIGIAT TITINAGIAGA GACGGGGITT CATCATITINA GICAGGCIGG TCICAGACIG CIGACCICAT
GATCCACACG CCIIGGCCIC CCAAAGIACI GGGATTACAG GCAIGAACCA CCACGCCCAT CIGATITCCC GITITCIGCA
GGGIAAAGIC TCAGGGCCGG CCCAITGNIT TCAGGANITI T

SEO ID NO:1334: (Length of Sequence = 267 Nucleotides)

MNATAACITI TIGITGAAAT TIAGAAAATG TGGATCITIT ATACTIGCIT TCCCTTTCT TCIGCCATCI TIATCITCIG CTGAAGGAGA CAAACAATAT TITAGGIGAC ATCIATCACI TIATGIAGGA CCIGCAAACA CTCATGTIGI CITCGGACAG ACAAATGGAG AATGIAAAATC TGITACACIG TGACAGGATA TAATINIGGA TIGCATAGGN TINCAACAAA GIGICIGIGI GATGANIAAA TGGIAAAATA TATTIAT

SEO ID NO:1335: (Length of Sequence = 279 Nucleotides)

GENTCITGIT AGAATGCAGA TTCTAATTAA AAATGTGTAG GACAGGGCCT GAGACTCGGT ATTTCTAACA AGTTCCCAAG
TGATACTAAT GCTACTGCTT CACAGATCAC ACTITAAATA GTAAGGTTCT TGAGAGAGAGT TAGTCTCAAG AGAAAAGAGA
CAAAAATCTC CAGAGCAGGA AGACCAAGAA AAAAAAATGG AAAGTAGCCA GTCGATTATC AACTAGATGG CCTTAGTGAG
ATTCTGCACA ATATTTCATC ATACAAAACT GNITTCCCA

SEO ID NO:1336: (Length of Sequence = 398 Nucleotides)

TTTTTTAAGC ACTCTTGTGT GGACTGGTCA AAGATGTTCC TAAAACAACA TTGCTGTCAC CAAGCCTCCC ATGANTTAGG CAGGCTCCTC CCATGTGGAT ACCTGCTTC) GCATAGTTGA TGAAGAGGAA GCATCCTCAG TCAAGCCTCC CAAGCAACAT ACCCTTAGGA AACCCCGCTG GTACCTGGCC TGINITTTGT AAGTATACAT CAGGCCAGGG GGCTGCTTGC CAAGCAACAT CATTGACTGC ATACTGTTTA GTGCATGCAT TACCAGGGCT CAAACATCCA AGTGATGCTA CCTGAATAAG TCGAGGAATT TITGATAATA AACATAAGCC AAATCCAAAA AAATGITCTG GGITTTTCCA TCATTTCCAC TCATTAGINC CAGGAAAA

SEO ID NO:1337: (Length of Sequence = 272 Nucleotides)
CTTTCCTCAG TATCACAGGT ACCIGITTIN CIGGAATITA TITAAAATGT CACCITGTAG TGITCCCTCT CTAGGGCTGT

TIGITICATI TOCCTOTGAA TGAATGCTGC CACACGGTCA TATGTGAGCC AAGTTTACAA GAATGGAGTT GCTGCTGAAG
AGATCTCTCA TTCATCTCCC CCAGTGCCTG TCCTTCACAA TCATAACGTT ACCCTTGCTT GACAAATATA CTGTATGGCA
AGTCATAAAG GTCTTWGAAC AGGACTTGAC CC

SEO ID NO:1338: (Length of Sequence = 212 Nucleotides)

TAGICCCCTT TATATAATAT AATCAAGTTC CICCATCIGG GCATICAGIT AAATTCIACA ACATTGCCAA AATCIGATTT GACICIACAG AATATGIATA GITTATTTAA CCAGATAGTA ATTTAAAAATT TTACAACATG CGTATTTCAT GTAATATTAA TAACAGTAAT TTAAAATTAAT ATTCAATACA TACCGTTTGA ATTTTTATAA GG

SEO ID NO:1339: (Length of Sequence = 280 Nucleotides)

TTTTTAGGAA TAACAAATGT TTATTCAGAA ATGGATAAGT AATACATAAT CACTCTTCAT CTCTTAATGC CCCTTCCTCT
CCTTCTGCAC AGGAGACACA GATGGGTAAC ATAGAGGCAT GGGAAGTGGA GGAGGACACA GGACTAGCCC ACCACCTTCT
CCTCCCGGTC TCCCAAGATG ACTGCTTATA GAGTGGNGGA GGCAAACAGG TCCCCTCAAT GTACCAGNTG GTCACCTATA
GCACCAGCTC CAGATGGCCA CGTGGCTGCA GCTGTACTCA

SEO ID NO:1340: (Length of Sequence = 324 Nucleotides)

CIGITOCACC TCAGATCATC AGGCATTAGA TTCTCATAAG GAGTGTGCAA CCTAGATCCC TCCCATGTGC TGTTCATAGC AGGATTTGCA CTCCTATAAG AATCTAATGC CACTGCAGAT CTGGCAGGAG GCGGAGCTGA TGGTGGGAAG GTGGTATTGC TCGCCCCCC GCCTACTGCT CACCTCCTGC TGTGGGGTCC AGTTCCCACC ACAGACCACT GGTCTNTGAC TCAGGGACCAC CTAC CTC AACANGGNTG AGGAAAACAA CTGGGTTCAT CACACAATTA TTTTAAAGTT CAGGTTTTNC AAATAACTTA

SEO ID NO:1341: (Length of Sequence = 376 Nucleotides)

CTAATCAAGG GTACAAGATG TCTAANTCAA AGGCCCAGCT CTGCCTACAA GTCAAATATC TAGGCCTAAT CTTGGCCAGA
GGAACCAGGG CCCTCAGCAA GGAATGANTA CAGCCTATAC TGGCTTATCC TCGCCCTAAG ACATTAAAAC AGTTGTGGGG
GTTCCTTGGA ATCACTGGCT TTTGCCGACT ATGGNTCCCC AGATACAGCG AGATACACTC TAAGGAGACC CAGAGGGCAA
ATACTCATCT AGTAGAATGG AGACCCAGAG GGCAAATACT CATCTAGTAG AATGGGGACC AGAGGCAGAA ACAGCCTTTC
AAAACCTTTA AAGCAGGCCC TTCTTNCAAG CTCCAGCCTT TAAGCCTTNC CACAGG

SEO ID NO:1342: (Length of Sequence = 335 Nucleotides)

ACCCTICCCC ACTCCCTGGT CCCCGGGAGC AGCTCCTTCT GCCCGANTNA CTCACAGTGC AGGGAAAGGA GGCAGGGAAA AGACCAGGAT TCTGTGAGTT CTGAGGGTTGC CACACACAAA GAAGCTGTGG TTTCTCTGCC TCGGCCACTG ATGAGACTAA AACTGGCTTC CCCTTGGAGA CGGCAGATTT CAGGCTGATC CCTGCTTAAG CCCTCTCATC CCCACGCTGG TCCTGGTATT GATACAAGAC CCAGCTGTG ACNAAGCCTC CAATCCTGGG GGTCCACGGA GCCTGGGGCT GANATTTCCA GGAACTATAC GCCAGTGGGC GCCCA

SEQ ID NO:1343: (Length of Sequence = 379 Nucleotides)

GAACCCAGGA GGCGGAGGTT GTAGTGAGCC AAGATCGTGC CATTGCACTC CAGCCTCGGC AACAAGAGCG AAACTCCATC
TTAAGAAAAA AAGAAGGGTG TGATAGTTAA ATTTATGCAT CAACTTGGCT AGGCAATGGT GTCCAGATAG TTGGTCAAAC
ATTATTCTAG ATGTTTCTGT GGAGGTTATT TTTTAGATGA GATTAGCCTT GTAAACTGGT GAAAATTGGG TGAAGGAGAT
TACCCTGCAT AGTGTGGTGG GTCTCATTTA ATCAGCTGGA GGCCTCAATA GGGAAAAAGA CTCACCCTNC CCTGGAGCAA
GAAGGAAATT CTTGCCCAGC AGAACTTCTT NGGGCAGCAG AATGCAACCA TAAACTCTT

SEO ID NO:1344: (Length of Sequence = 400 Nucleotides)

GACGGATGGG ATCGGGGCTG TGCTCTGCAG GTCCTCCCCA GAGATGTTGT CATACTGCGA GGGATGCCGC TCGTAGGACA CCCTGCAGCC AGAGCCGTCC GCCGTCTGGAA AGAGCGCCGAA GAAATGGAGA CCCAGACTCC CCCAAAGACT CTGGCAACAG GCTAAGGTC CAGGGCCGTC TGCTGAGGTA TCTGGTCTGC GTTAGAGAGG TCTTNCTGGA GGAATTCATA GTCGGGATCA TAGCAGATCT TGTCCCCTTT CTATACCATC TGTCCTATTT GGAGATNGCT

SEQ ID NO:1345: (Length of Sequence = 347 Nucleotides)

SEQ ID NO:1346: (Length of Sequence = 287 Nucleotides)

CAAGICAATA CCCATAATTA AGICAAGTIG CCAGCCITAA TIATATIINI NICICGCICG TICACICIC CICICCITCC
CICCITCCCT CICIGCCCCA CCCCCGIGIA CATTATATAC CAATTCATIG GAGATATATA TATGINIGIN INIGNGINIG
TGIGIGINIC TGIGIGIGIG TGIGIGITIAA AGAAGCAGGA TGICITACAC AGATGITICA TATATTGAGG NATIACAGAG
TAATIACAGG GAAAGGIATI ACACTGITCT TCAACACCCT AGGCAGT

SEO ID NO:1347: (Length of Sequence = 295 Nucleotides)

ATTAAACAAC TITTITAAAC TITTIGIGCA CAGGACAGAA AACTGCCTGT ACATGCTATG TCCACTITTG GAACACAGAT TITTIAACAAT TATGAATGCA CAAAATCTTA CATATCATGC AACTCTATGC CAAGAACCCA ACTTTCTTCC ATGCAACAGA TATGAAGATC TAAATGGAAA CCTAGCTAAG TCTTAAACAC TITTCCAGTA GCAAGTATAA TATATGTTGT TGAGGGAAAA CCAGTCTTAA CAATTACTTG TACACAATAT TCATGTGCCA AATACAATCA CAGGA

SEO ID NO:1348: (Length of Sequence = 332 Nucleotides)

AGICCCIGCT ATGIGGATAT TIGGTAGCAA TGACTGATGT GGAAACTACA TATGCAGATT TIATTGCTTC AGGAAGAACA GGTAGAAGAA ATGCAATACA TGATATCCTG GTTTCCTCTG CAAGTGGCAA CAGCAATGAA TTAGCCTTGA AATTAGCAGG TCTTGATATC AACAAGACAG AAGGTGAAGA AGATGCACAA CGAANITCTA CAGAACAAAG TGGNGGAAGC CCAGGGAGAA GCAGCAAAAAT CTGAAAGCTT AACACCCCAC TTTGACCCTC GGCCACACCT GAAAATGTCT CAAATCTCCA GGGNGTATCT GGGAATGCAT TT

SEO ID NO:1349: (Length of Sequence = 296 Micleotides)

GCCCCAAAAA CAATGACACA AAATTCATTI GGITAATTCA TGTAAAGGAA AAAACAGCAA CACCACCACA CAAACAGGAA AGTGGGAGTA TGATTAGGAG GGGTGAGATG AAAACTATTI TACAGTAACA TTTCCACCAA AAGACTGTCC TAAGAACAGG CTGTCAATAC AGTTCACAGG GAAAAAGCAA ATGTGGTATT TTTTTGTATT TTTTAAAAGC TCCCTGGGTC CCAGGTGTTT TGCAGTTTTC AAGGNCTTAT CTGCTAAAGG AATGCCCTTT TAGGGTCACA GCAGGT

SEO ID NO:1350: (Length of Sequence = 317 Nucleotides)

CIGITGCCCA GGCTAGAATG CAATGNCGIG ATCITGGCIC TCACIGCAAC CICCACCICC CAGGITCAAG TGATICICCT GCCICANICI CCCIAGIAGC TGGGATIACA GGIGITCACC ACCACGCCAG GCTAATTITI GIATTITIAG TAGAGAAGGG GITTCACCAT GITGGCCAAC CICGAACTCC CAACCTCAGG TGATCCACCT GCCTCAGCCT CCCAAAGTGC TGGGATTACA GGCATGAGCC ACTGIGCCTG GGCCAATAAA CTATATTITIN TCAAGCCAAA GTAGGACAAG CACAGITTITI AAAAGGG

SEQ ID NO:1351: (Length of Sequence = 349 Nucleotides)

CGGATGGGTG GGATGAGACT TCAGCTTTAT TGGAAATGIT TTATTTCCTT ATCTAAAAAA ATACTAGAAA GAAATACAAC
AAAATGITAA CAGTTGTTAA TGTCGGCCTC TGTAAATATA GATATTGTGT TACTTTAGTC TTTTTTTTAA TCTCAACTAA
ATTAAAAAAAG GAATTTTAGT CTTTTTTTAT CTCAACTAAA TTAAAAAAGG AATTTTAAAA CCCTAGTGTT ACATGCAAGT
GAGTCCAATA ATGGCAAAAT AATAATGAGG NTACATAGGA AGGGTGACCT AAATTTTAAT GGGTGAATAC TGGGTCCCCG
GTACAAGTTT GANAAATTTT GAATTTCCG

SEO ID NO:1352: (Length of Sequence = 304 Nucleotides)

TTTTTATACT ATTTAAAAGA ATCCTTAAAT GATGGGTATT CTCTAAAGCA TGCGGGGCTT AAAACCTAGA TGATGGATTG
ATAGGTGCAG CAAACCACCG TGGCACATAT ATACCTATGT AACAAACCTG CACATTCAGC ACATGTATCC CAAGACTTAA
AGTAAAAGTA AAAATTAAAA AAGATGGGTA TTCTATATTT ATCTTTCATG TTACATTTTT CTTTGTGGGG TTTCTAAATA
AAACTTGTAA CATGAATGTT TTATTCTCAT TCTGTATTTT AAAAAGAAGC TGAGTAACAA AAGG

SEO ID NO:1353: (Length of Sequence = 307 Nucleotides)

CTTAGTCTGA CATTAGGTTA TGAGAAGTAC AAAAGATCCA CAAGTACAAA AAAATCTGTA TAGCTTTGCG GTAGTTGAAA
AAAATGCAAG AGAACAAAAA AATTTTTTGA GTAATATTCA TCTCTGCAGA TCTGAGTGAC AGTCCGCTTG AAACACCGCT
GTAAAAGTGG TAAAAAATGA TTTCATTGTG ATTATGTTAA AATTTTTTGAT GTCTCTINTTA CTTGTTTTTAG GGGAATCTGG
TCTTCCTGNC ATTTATACCT GGATANGTNC CTTTCCCTGT AATTTTTNCT GAAAGGCTCC AATTTCC

SEO ID NO:1354: (Length of Sequence = 407 Nucleotides)

GIGAAGITAA GCAGCAAGGG CIGAGAACCG CIGCICCAGA GAGGCCAGGA GGICIGGICA GAGGCIGGGG CCCCAGCCCC
CAGGCACCIC ICIGIGICAG ITICCCIGGA GAAGICATGA GITIGAAGAG TAGGCAGAGG CCAGGIGICA TCACIGAGIC
ACICATCAAT GGCCAATGAG AGINCAAAGG GIAGCICCIGA GCACAGGATG INIAGCAAGA CICCIGGGII CAGCICCCAG
TCCCCACCANI GCCAAGIGGG GGATCCITAG CAAGGIACTI ACCITITINN TGCCICIGII TCTACGGCIG CAAAATGGGC
ACAATAATGI CAGATICATG AGGGATAATG AGGACTAAAA TTAGGNIAAT INCCIATAAG CIGCITCIAA ACGIATITAC
TITATAAA

SEO ID NO:1355: (Length of Sequence = 355 Nucleotides)

ATTACTATTT GCCTCTATAG GAGGITTCAT TAGGCATCIN CITCATTATG AGGGCAATAT AATCAAACAC TTATCAGTAC AAGGCAGAGA GACCGGGACT AGCTGCCTAC ACATCCTCAA TGAGCTTTAG GAAATTTGAA GGAAACATGG ACTGAAAAATC TTCTGGTGGC AGGTACTCTC ATGTGTTGTC CTATCTGATG CTCTCAACAA CCTCTAGGGG TAGATATTGT GACCCTCATC

TTGCAGAGC CTTGGCTCAA GTATATGCTC AGAATCACAG AGCTGGAAGA TAAACTTGGG TCTCTCTAGT GCCAGAGNCC ATGNCCTCTG ATCTCTCAAG GGCAGAGGTA TTACC

SEO ID NO:1356: (Length of Sequence = 406 Nucleotides)

TITITITIAG TIATTCACT CICTCIGITA AATITATCIG ATAGGATICI GCAGAGAACA AAATICAACA GGGCCCIGIG GAGCAAGGAG CCCCTITICC CIATCICCT CCICTAAGAG CIACACCCAG ACCAGCIGI TATCAGCGGA GGCCCCGCIG CICCCCATGA GAACGCIGGI GGAAGACGAA GGIGATGGCA GTIGAAGGCAG CATCCCAGGC AGCCIGGAGI ACCICCATCCC GGAGCCCCCA CITATCAGIG CAGIGGITCC ACCCIGCCAG GGICINAAGI GCAGICAGAA CCATCAGGGG GINGCCGGAT CIGACGGCIG TINACACAAC GIOGGCAGIG CAAACCTAGG GACAGAAGGC ACANCINAAG TCACINCAGA TCCCATCITC CIACIG

SEO ID NO:1357: (Length of Sequence = 231 Nucleotides)

TTCACAAAG AATTTATGAT TGCTTCACCA GGTCACTAGT GAGCTAAAGT CAAGGAATGA CTACAATCTT GTAGCATTTT
AAAGTGATTA GAATTTGAGA AACTTTTACT ACATTATGTG TTACTATCAT AAGAACACTC CTTTGGGGGC ATTTGAATAA
TAAAAAGGNC TACATTCTTT GCACCANGTG NTCATTTTCA CCCACATTCC AGTATTTTINC TCTAACTTGG G

SEO ID NO:1358: (Length of Sequence = 302 Nucleotides)

CACAACTAAT TIGTAAGCCC CTTGAGCGCA GGAACTGGTT TITTAAAGAA TGATGTATIC TICACAGTGC TITCCCTTTC
TGTTACCCAG GGAGCACATG GCAATATAAG GGCTCCTGGG ATTGANTCTT AAGTACAGAG AAAACCTAAG AATNCTTTTA
GATAGACACA TAAGAGACCA CNAGAGAAGA GCAGATTCTA AGGTATNTGT GAGAAACGTT ATGTAATGAA AAGATAATTG
ATGACACACA CTTCCAGAGN GTGCTGGCGA GATTTGATTC AAAAGCACAC GGCTAGGGCA CT

SEO ID NO:1359: (Length of Sequence = 356 Nucleotides)

TAATGATGAG CCTCTGGGIG CAGGAGAGG ATAGGACTIG ATGCTTTCCA GGGGAAATAT TIAAAATTIC AGIACTAAGT
TAAGICTGIA TCATTTTACT TTTTTTATAG TTTCTTATTT TATGITGIAT GAGATGAAA GCTTGCACAT AAAAGATGAT
AAGAAATTAG AATTCATCGI TTCTGTTGIA CCAAGAAGAA CCTTAGTGAT CTCTAAAAGA ATTGTTGITA AAATATGGAT
TCTNCTTTCC TTCTAGTACT CCCCTAGCAT GACANTGAGC GTGTGATCCA TTACCAAGTC TCCTCATGAA AACCACAGTG
AGTCAGCCCT TCACAGAACT ACTACGGAG GAAATT

SEO ID NO:1360: (Length of Sequence = 366 Nucleotides)

SEO ID NO:1361: (Length of Sequence = 347 Mucleotides)

CCTCCTACTG TCTTGTCTGT GGGACAGTTG CCTCCCCCTC ATCTCCAGTG ACTCAGCCTA CACAAGGGAG GACCAACAGG
NTCTAGTTTT TCCACGTGAT GGAGTTCCAA GCTTTTTTTT TTGATYTGTT TTGTTTCGCA AAATAAAAAC AATACACATT
CCAAGAGAAA TGAATGCATC TNITGACACG TCTCTATTTTC TCATTTACAT ATGTACACAC GNCCCTTGAG TCGCTGCTCT
TGACACGGCC CNGTGTGGAC GGGTCAGGCC CGAGGCCCCT CGGGAGCAGA CCTGTAGCTC TCTGGGGGAT CAGGGCTTCC
ATTAGGGAGA AAGTATTAGC AGTTTCT

SEO ID NO:1362: (Length of Sequence = 358 Nucleotides)

CCATTCATTC ATTCATTCAA CAATATTCAT TCAACAAATG AAGCAAAGGA GCACACAGCC AAGTGATGGA GCAAAATCAC
AAATTAAAAG GTAATTCAGG CCAGGTGAGG TGGCTCATGC CTGTAATCCC AGCACTTTGG GAGGCCGAGG CAGGTGGACC
AGCTGAGGCC AGGAGTTTNA GACCAGCCTG GCCAACATGA TGCAACCCCG TNINTACTGA AAATACAAAA ACAAACAAAC
AACATAAAAA AATTAGCCAG GTATNGTTTG CAGGCGNCTG TAATTNCAGC TTAGTCAGGA GGCTTTGGCA NGGGCTTCAG
TTAGCCAAGA TCGGACCCTT NCACTTTCAG CCTGGGTA

SEO ID NO:1363: (Length of Sequence = 312 Nucleotides)

TATTIAAATA ACGIGCAATT TCATAAATCA GCACATITAC TAGATAGGIA GGATACITIT NATCCATITG TGIGITAAAA
AATTAGCGCA TGITTCTCTT TATGCCCACT TGIATTAGCA GAATAGIGIT TICGGATTCC CIGAATGGIT CIGIATIGAG
TCTGIATAGA CCCCGAAGGA AAAGGAGGAA TICGCCGIGC CCGAGAATAG CICCGICCAG CAGITTANGG NAGAAATCTC
TAAACGITTT AAATCACATA CIGACCAACI TGIGITGATA TITGCIGGAA AAATTITGAA AGNICAAGAT AC

SEO ID NO:1364: (Length of Sequence = 345 Nucleotides)

CTGACAGATT TACAGATGCT GACCTATTGA AAAATACCAC AGCCAGAATG GGCTAAACAG GTATATAGTT AATACAACCA CCACCATCCT TTACTITTAA CATAGCTCTT AGTAGGAATT TCATAAAANT GGACATCACA GCTAAAATGC ATTATTAATT CTCCTATCTG CTGACAATAA AAAAGCAGCA AACTCACATGA TTTCTATTTA AATGCACTAG ATGGGAATAT CATGITCTAG GGGTGTTTGC CTTCAAACCA AACCCACAGC AACACACACA AGCAATTTCG GTATCCACCA TTTTAAATTC ACAATCTGAG NCTAAATGAA TGGCTATTTA TATTT

SEQ ID NO:1365: (Length of Sequence = 255 Nucleotides)

CTCCAGAAAG CCATTGATCT GGTGACGAAA GCCACAGAGG AGGACAAAGC CAAGANCTAC GAGGAGGCGC TGCGGCTGTA
CCAGCATGCG GTGGAGTACT TCCTCCACGC TATCAAGTAT GAGGCCCACA GCGACAAGGC CAAGGAGAGC ATTCGAGCCA
AGTGCGTGCA GTACCTAGAC CGGGCCGAGA AGCTGAAGGA TTATTTACGA AGCAAAGAGA AACACGGCAA GAAGCCAGTC
AAAGAGAACC AGAGT

SEO ID NO:1366: (Length of Sequence = 322 Nucleotides)

AAAAAAAAA TICCAAAGAA ACAGAGTAAT TITCCTCCTT GCCTCAGCCC TAAGTCATCT CCCAGACAAA AAAGCAATCA
TCATTGTCAA ATTTAAAAGG GAAAAGGAAA GACTITTATT TGANTGAAAA GATTITITTC AGTGTGATAG AGAGGGAAGA
CTGAAAATAAA CAGAATTTAC AACCITCGCA CCTTTGCACC TTCCTCTTCT AGCAGTATGG CAAACTAAAT AACTTGCACT
GAAAACGGGT TAAAAAAGCTG TATACITTTT TAAAAAAATAT ATTIMENITA TGTCATTGAT CTGCACAGTT TTGAATACAA
AA

SEO ID NO:1367: (Length of Sequence = 349 Nucleotides)

GAAAACAAGG TCAACATCAC TCATCATTAG AGAAATGCAA ATCAAAACCA TAGTGAAATA CCATCTCACA CCAGTCAGAA
TGGCTGCTAC TAGAAATAAC ATGCTGGTGA GGCTGCAGAG AGAAAGGAAT GTTTATACAC TGTTAATGGG AGINTAATTA
GTTCAACCAT TGTGGTAGAC AGTGTGAAAA TTCCTCAAAG ACCTAGAGAC AGAAATACCA TTTGACTGAG CAATCTCATT
ACTGGGTATA TAGCCAAAGG AATATAAATT: GTTCTACTGT AGAGAAAACA TGCATGCATG TTTGTTTGCA GCACTATTTC
ACAAGAGCAA ACACATGGGA TCAACTTAA

SEO ID NO:1368: (Length of Sequence = 379 Nucleotides)

CIGGGACAGA GACCITIGCA TIGCICCAIG TGITGGCITC AGCIGGGACA GAGACCITIG CATIGCICCA TGIGITGGGG
CAGGICTICC ATTICAATCI CCICTGCCCI AATTIATIAG CCATACITGI GCIATITATT ACTITIAAAC CCIAATCCIT
TITCCGIAAT TIGITIACAT TITGCAGAGI GCCAGCATTI TACAATGIGI CITTTATGIC TCACAGAGGI CATCATTAAG
TTAGACCITT GGCITCATGI GICICCCGAG AGATGGITTA TAAAATTIGC AINCITCIGG CACAGGIGGI GIGGCITAGG
GATTAGGACA CAGCCIGCCI GAGITCACAC CICATCICCC CCACTIAACA CIGATAATT

SEO ID NO:1369: (Length of Sequence = 319 Nucleotides)

ATTTCIGGIC TAAGITITAT TATTTCCTTT CITCIGCITG TITTAGGCIG ATATTGCACT TCTTACTCCA GITTTCIAAG GIGGAAGCIT CGACIATIGA TITCAAATCI TITINCITIN CIAATCIATG CATTCAATGI TATAAGITIC TGIGAAGCAG TGATTTCATT GCATCCCACA TITTGATAGG TIATATTTCC ATTTAGITAC AAATAATTTA AATTTCCCTT GAGATTTCTG CTITGACITA TGIGITATTT GGAAGIGIAT TITTATTCTC CAAATATTTA GAGATTTGCA GCIGICITTA TGITATTAA

SEO ID NO:1370: (Length of Sequence = 343 Nucleotides)

GGAAAACATA AATNITGACA AGTAGITCAA GACIGITGGG ATAAACITAG CIAGAGIGCA GGICATAACI ACCCATCITI
ATAAGGAAGC TGAAAAGGGA AGTATGAGGA CAGGGAGAAC AATGACITIN TCICTCAAGC TIGACITAAA CCACCAGGAA
AGTICTTAAAA GCCAAAGCCT TICTCAGACT CICACCAAAC CATAAGAGIC AGAAAAATGG TCGITITCAA AGGAGTAGAA
AATTCIGIAC AAAGTAAACA ACAGCTGAAG CAGGAAAGGN ACATACATIT NNTCACTIAG TGGCACGCAG GCAAAACAGA
ACATAGGGCC AGCTTGGITA TIT

SEO ID NO:1371: (Length of Sequence = 295 Nucleotides)

SEO ID NO:1372: (Length of Sequence = 340 Nucleotides)

TITGCTITCA GATAATGITI CIGIATACIT TATAAATGCT ATCIGIGGIA TCTCCIGIAT AATINACAAT GITTGCATGI
AAAAAACAAA ACCCATAGAC CITAAAAAAA AGAAAAAAAG AAATATACAC TATACATAGG CACAGCITAT GCCCAGAGCA
TAGCAGGIGC ATAAAACACT GITGCTATAA ATGCAAGAAA AAGGICATIT AACCACAATC ACATTITITI NCATAAGAGN
GTCIGAAATC TATACAATAT ATACATCTAT GITTCAATGI GGAAATAATA TTCTTTTAAA TTTCAAGGCG TGTTATACCC
CTGCAGGCCT GCATAAATGG

. <u>SEQ ID NO:1373:</u> (Length of Sequence = 315 Nucleotides)

AATOCTGGGG GTGATTTAGA ACTTAGAGGC ATTCTCAAAA TGGACCAAGC TAAATGGTAG CCITTATTIN CTGTAATGAT
TCACCATGGG AAAATTAGTA ATTCTTTAAA CTTCTTACTT AATCTTATAT GTATTCCAAA TTINCTAAAA AGAAATTAAC
CTAGAGGTTT TACAGAACTC CATTTTTTT TTATTTNCCA GAAAGGAAAA ATTTATCTGT NCTGTNATTT TGTTAAAAAT
CCTATTCCAG CTACTACTAT GGAAAAAGGA AAAGAAGAAA GGAGGAAAGG AAGGGAGAG GGAAAGGAAG GGACG

SEO ID NO:1374: (Length of Sequence = 327 Mucleotides)

GAGCCAGTGG TGGCCCCCAA CAGCCCAATC TGGTACTCAG TCCAGCCTAT CAGCAGAGAG CAGATGGGAC AAATGCTGAC GCGGATCCTG GTGATAAGAG AAATTCAGGA GGCCATCGCA GTGGCCAATG CAAGCACTAT GCACTGAGAT GCCTTGGCCA AAAGGAAAAT ATAAAAGAAA ATAAAATCTC ACATTGGTGC TTAGCAGGAG AATTTTTAAA GACTTACAAA TCAACAAGCT GTTCAAATAA ATAATGAATG CTGCAGCTGG CTCTTACATG GGGCTTTNAG TGTCCCANTA GTAGCAGATG TCCCAGTTCT ATAAAAT

SEO ID NO:1375: (Length of Sequence = 338 Nucleotides)

TGCATGGAAA CITAATCTAT TCAGGTCCCA ACTITCAGGC TTCTCTGCTC TGACAAGTAC TAGAGGCCAA TATGATAGAC
TAGTCTGAGT TGGATGCAAG TTAGCCATTT CCAGGAATGA TACCAGGATA AGTATAATGG TCGTGAATAT AACCGGATTT
TAAGGGAGAA TGATTACACC TGGAAACAAA CTGTCAATAC ACAAGTAACT AGTTGTTAAA GATTTCTAAT TTTGACCAAA
GATTTTTACT TTCCTGGTAT AGAAATGGAA ATAAACATTN ACACTTTAGG TTTTGAAAGC AACCACCTCC TAACACGGTT
CTGAGTTGGG GGCCAACA

SEQ ID NO:1376: (Length of Sequence = 307 Nucleotides)

CAAGCCTCCC TCAAAAAAAT CCCCAGAGTA ATGAAAATAC AAAGTCTGCT TGTTCAAAAT TATGGTGCGA ATAAAAAAGG AAAGGGAGGA AGTGATGGAG TAAAGTTCAG ATTAAAAATA AACGGAAAGT CACAACAGTC GAAAGGTGGA AAAAAACCGC AAATGCCCAT GANCIGATGA ATGGATAAAC AAAATTTGGT GTGTGTGTAT ATATGTGTAC AAACTTCCTT TTTATGATGA AATAGTATTT CATTGTGTGT GCACATGTTN CACACACANT TTAAATAGTA TTTCGTCATA AAAAAAG

SEQ ID NO:1377: (Length of Sequence = 353 Nucleotides)

TGGAATACAC TIGIGAATAC AGIGIGTAGG ATACATTAAC AGITTICIGA GIGGGCIGCT CITTITICCT CAATACIGTA
TATATITINN TIAAGCICIT CITTAAAAGA TAAATATITI TCATACITCT CITAAATCCT CAAGGATTAA CICIGAGICA
CCATTIGIGG TATITTAAAT CCITTTAAAT AAATCICIGI ATTIGCAACT GCATCAAAAC AGIAAAACAT TICACAGGGT
AGGATCIGAT GACCATITTA TAATCAACAT TITTAGGTAC CACAAGAL "ACTITATGAG CATCCACTGA AATTATGGGC
ATTATGICAT ATAAATATCC AAAAATCCAT TIT

SEO ID NO:1378: (Length of Sequence = 315 Nucleotides)

GATTTGGCAA ATATTTGGGT GAGATTTGAA AATAAATTAC ACCACTGCTG CACAAGTTAA TGTGAATCAA GCATCTGTTT
ATTTCATTCA GTTTATGCCT TTTTTCTTT TTTTGTGCAG TGCAGTTGGG GTCACAGACT CTCAATTTGA CAAGACACTT
TAAAAGCAGG AGTAGAAATT AGGCCCGGGT TTTACAACTA TTACAAGGAAC TGTCATAACA AACTTCAAGT GGATCAGTTT
ATTTCTGATT TAACTTGGGG ATAAACAGTG TTCAATATTT TCCAAAAAGAT TCTCCCCATA TAGAAGTCCC AAAAG

SEO ID NO:1379: (Length of Sequence = 352 Nucleotides)

ACCECAAAAT TIAGCIGITI ATTAGGITGC AAGICTCICC TICTCICCCI GCITTCICIT TCINCITITI CICCCCACAA ATCCCCCCAA AACACATACA AAAAAGAGAAA ACTAGAAGCA AGATTGGGIC AAACATGAAG AACACAGAAA GCNTATTAAA TAGCTAGCIT TAAAAGGGCIC TITTICAGIT TGAACAAAAG TAAAACGITC TCAAAAAGCAA AAACAGAAAA CAGAGCITCC ACCCAGATTG TGCAACTTAA TGAGAGGAGG TTAGTGCTGA TAAACCCATT GTGAAATCTA TTATAAAGTG ACAGGITTTT CAAGCAAGGA AATCCAATCC AGTTGGGGGT TG

SEO ID NO:1380: (Length of Sequence = 261 Nucleotides)

AAAAATTIAL TCAAGACGIG AATAGATATI NCIGCAAAGA AAACATACAA GIGGICAATA GITATATAAA AGGTATTCAA
TATCACTAAT CATCAGGGAA AIGCAAATCA AAACCACAAT GAGTTATCIN CICATACCIT INAIGMIGGC TAATATTAAN
CGAGAGATAA CAASTGITTA TGGGGGIGIG GNGAAAAGAG AATGITCGAA CACICTIGGI TGAAATATAA GITGGIAGAA
CCATTATGCA AAACAGTATG A

SEO ID NO:1381: (Length of Sequence = 273 Nucleotides)

GCCACTACAC TCCAGCCTGG GACACAGAGC AAGACTCCCT CCCAAAAAAA AAAAAAAAA TTATTAGAAA GAGGAAGAGA GAGATGACAA AGCCTTTTAC AGTTGGGTGT TGGGAGTTAG AGACCCAGTA CCCCAGCCTG ACATACCTAC AGAAGCAGTG AATTTACTTA TTTACTGTTA TGAAAAAAAAT AGATGCTGCC AGCCGTGCAC AGCAGAAACT ACTATTGANT CATATGGTTT TAGCCTTCAC CTTTAAATAT GTCTAATTAT ATG

SEO ID NO:1382: (Length of Sequence = 296 Nucleotides)

CTCCACAGCT GCCACATAGA ACAAGCAAAT CTGACATCAC AGCTCTTTTA AAAATCTCCC AGAATTCTAC ACTGGAATAA
AGATCACCCA GTAAACTCAG CTATGTTGAT TCGTAGGAAT TTCTCCTTGG AGTTAATAAT AATCATTAGA AAAAAAATAC
AGGAAGAAAT AACTTCCTCC TATTCTTATT GTGATAAATT GTAACAATAG CAGACATTCG TATATAGATC CTATAAGCGA
CAAGAGGGAA AATAGGATTT GCAANITAAG CATCTGGAAT AAATATTTTTA GGAAAA

SEO ID NO:1383: (Length of Sequence = 293 Nucleotides)

CCAAGGACCG GGCCGTGCGG CTGCTCTGGG ACCGCTTCGT GCGGGGGTGC CGCGCCGACT GGTACGGAGG CAATNACCGC TCGGTCATCT GCTCTGACCA CTTTNCCCCA GCCTGTTTTN ACGTCTCTTC GGTTATCCAG AAGAACCTGC GCTTCTCCCA GCGNCTGAGG CTGGTGGCAG GCGCCGTGCC CACCCTGCAN CNGGTGCCCG CCCCGGCACC TAAGAGGGGA GAGGAGGGAG ACCAAGCAGG NCGCTGGAC ACGAGAGGAG AGCTTCAGGC AGCCAGGNAT TCT

SEO ID NO:1384: (Length of Sequence = 378 Nucleotides)

GGTGGTTTTG ACATGIAGAA AATAAGATGG AAGGCTGAAC TAGGGCAGTG GTGTTGGCAA ATAATCAGAT TICAGGAATA
TCACAAAGTG AGGNGCCCAG GATTCATGAC CATTTINATG TAGGAATAAG GGAGGAGCCT AGGATGACTC CCCCAAGTTT
CTGGCTCGAG TAACTGGGAT ATCAACAAGT CATTTAGCAA AATAGAGAAA ATAGGAGAAG CAGCAATTTG AGATAGAGAT
AGAGGCCAATA TAAAGNNTTA TATATTGACC ATGGTAAATC ACCTAAATTC AGAAAGTTGT AGAAAACTTG GGTCTGGANC
TCAGGAAAGA CACTGGATAT GTAGATTTGG AAAGTTATCA ATCTCAAAGT GATTGCTT

SEO ID NO:1385: (Length of Sequence = 204 Nucleotides)

TCATTCTTGG GTGTTTCTCG CAGAGGAGGG NTTTGGCAGG GTCATAGGAC AATAGTGGAG GGAAGGTCAG CAGACAAACA AGTGAACAAA GGTCTCTGGT TTTCNTAGGC AGAGGACCCC GAGGCCTTCC GCAGTGTTTG TTTCCCTGGG TACTTNAGAT TAGGGAGTGG TGATGACTCT TAACGAGCAT GCTGCCTTCA AGCA

SEO ID NO:1386: (Length of Sequence = 238 Nucleotides)

CCCCATCATG GGCAGCCAGA GCTCCAAGGC TCCCCGGGGC GACGTGACCG CCGAGGAGGC AGCAGGCGCT TCCCCCGCGA AGGCCAACGG CATGGAGAAT GGCCACGTGA AAAGCAATGG AGACTTATCC CCCAAGGGTG AAGGGAGGTC GCCCCTGTN AACGGAACAN ATGAGGCAGC CGGGGCCACT GGCGATGCCA TCCAGGCCAGC ACCCCCTAGC CAGGGTGCTG AGGCCAAG

SEO ID NO:1387: (Length of Sequence = 295 Nucleotides)

TITITITITI TITITITITI TITIANITAG GCAAGAAGA GTGTGAGTAA TIGAGGAAAA ACTGACAGAT GCTTTTCCTA
ATACCAAAAT TGAGCTTACA ATTAGGAACT GAGTATGTGT AACAGGNTAC AGGTGACAGT GAAGATAGAA GAACCACGNT
GACCACAGAC TCAATGTGCT CTGTAACATC GCACAGTTTA CCCAGCATGA CTTTCCTTAG GAGGCCCCCT CCTCACGCTA
GAGTAAAAGT CCCAGTTAAG TGAAGCCTAC CAGAAGAACT AGTALLAGAA GTTT

SEO ID NO:1388: (Length of Sequence = 201 Nucleotides)

GCTAGINATC TCTCAGACAC TTGGTCGGTA GAAAAGATCC CGCACCATCC TCCAGGNTCC AATGGCCTTG GAGAGAGGGC
TGCAGGGCCC ACGGNCATTG CTGACTCTTT AGAACGTGCT GACATGGAGC CAGACCACTC GGCCCTGAGT GCGGCGAGGA
CCCTNTTINT GGATGTGGAG GAGCGCGGC CGGAGCATTG T

SEO ID NO:1389: (Length of Sequence = 399 Nucleotides)

GGTGCCCTGT TATCIGGTAA AAGAGCCACT TATGACCTCA GGTGCTACTT AACCTGGGGG GCAATTGTTT CTTAGGCCTA
GCAGATGTTT GGGATGACAC TAAAAACTCA GTGGTGAGAT GATTCCCTTA GCAAGATTGC TGAAGTTAGG TTTAGACGTG
GGAGGGTGGG TATGTGAGACA ATGGTGCCAA TAGCGGCTCT TTATTTGCCT TGTCCTCATT ACTGCCATCA GGAAGGTGCT
ACTGGCCTCG AGCCAGGGTG TTCATAATCT GGCCTTGGGT TAACCAGACA AATAGAACTT CTTTTCCTAG ACTGTTGGCT
TINIGGAGGT TGGCAGCCTC TATCACAGGN TAAAATTTCC CAAATCCATT TACCCAGTAT ATTCACTACA ATTTTTTTCC

SEO ID NO:1390: (Length of Sequence = 381 Nucleotides)

GGATTGAGGT GAAGATACAA CAAAGAAAGG AAATTGAACG GAATTATTAA GAGGGTCAAG TTTGATAGGC AGATAAGACT
AGGTATCAGC AAGACATTC AAACAAAAGG AACATTATGT AATTTTTTAA AAAAATACAT GAAAATAATA TTTAANCAAG
GAAGGAATAT GATAAAAGAN GGATAGTTAG TAAAATTTGG ATAACATAAA GATTATTGAA TCTCCAGTCG TCAAATTTAT
CCTAAACTAC TGGGGAGAGG TCTCATGTCA GATTTTGATT ATCGAGAAAG AGGGGTCAAG AGTATAACNG AAATTCCTTT
TTGTTTTGAA CTTCCAGTGT CCCNCTATTG TGGGCAAATA TCAAATTCAA ACCAAATATA C

SEO ID NO:1391: (Length of Sequence = 327 Nucleotides)

GAAGAAGTCC TTCTTAAGCA AGGCTTACAG ACTCCCAGGG AGAACAAAAT CTCTTTATCT CTCTGGGGTT TTAGGACCCT CATCAAGTCA TAGAATTGAA ATAGAGAACA TCAATTGINC AACTTTTTAA TTTTAATAGT TTTTGTAGTA CATAAAAAATC ATGTTATGAA TTATTTTGTA GTTTTAATTA TAACTTTTTT AGCACTTTTA CCATATTCTT AAAAAATTAAA AATTATGAGT NCTGAGAAAAG CAGTGAAATC ACATATAGGT ATTTGATTAA CTTTTATGTG ATCTTTACC TCAAGCTAAT GTTTCTTAAA ATCAAGG

SEO ID NO:1392: (Length of Sequence = 223 Nucleotides)

TTTTTTAATA TTTAAAACAA TTTTATTCAT GAAAATATGC TGTACAATGC ACTCTACACA GCCTCGACAC GGCACACACG CACACGCACA CTCTGACGGC ACGGCCACGG TACACTGCCT ACGATACGCG CCGGGGACGC CGCGCCCACC GCCCGTCCCG GCCGGACACT TATAAATATG GGAGAAGGGC CAGAACTNGC GCGGAGAAAG GGGCGTCGGG GTT

SEO ID NO:1393: (Length of Sequence = 296 Nucleotides)

GAAAGITTAT TATITCCCAA TGINCITTAC ATTINCATIT GGAAATATCA TICCIGACAG AAATAGNTAC ATTATACCTT
CGAAAGCAGA AAGATCITAA TIAATTAAAA CAGITTACAT TIACCITTAGC ATTAGGTCTG GCTGGCTAAT TICAAAGGAT
TAAAAAATTGC ACCNATTTGG GCCAACTGGG GTCCTGAATA ATTATCCNGG GTAAAAGTAT AATATTTCAT ACTITTATACA
TITTGCTTCA TCACACATTT ACTITCCACA CAGTGNTCAA CTTCACACTTT AAAAAG

SEO ID NO:1394: (Length of Sequence = 281 Nucleotides)

SEO ID NO:1395: (Length of Sequence = 323 Nucleotides)

- CTTTTTTAA GATTTCAAAC TGGGTTACAC ACTGGAAAAG GCTGGGTTAA GGGCCGAAAT TTAATAAATC TGTACTGATA
- ACTARAGECT ACAGAGATTI CATATATTIT TITTAACTIT TAGARATCAG AGTECTIATA ARATEGCTEG CICATEGCTE TETCACCCAG CATCICTEAC GCCGCCTCCT AGCCTTCGTT GGTGAGATAA CCNGGNATAG TGATTCCATE CGTARACAAC
- AGAATACTA AACCAATAAA ACTAGCTTAT CATGCAAATA TTANGGCATC TAGAAAGTCA GTTAAAATAA TATTGTCATA

GAG

SEO ID NO:1396: (Length of Sequence = 384 Nucleotides)

TECCTCCCEG GITCATECGA TICTNCCECC TCAGTCTCIT GAGTAGCTGG GATTACAGGC ATGCACCACC ATGCCTAATT
TTTGTACTGA TGCCAGCACT TTCTTAGCAA CCCCAGCTGG TGTCCTAGTA TGCCCCCTCC AGTCCACTGT CTCTGGGCCC
AGTTCAGCGC TAGGACTTGC TTAAGAGTTT CAGTCCTTGT AGCCTATACT GCCTTTNACG TTTATTTAGA GATCTAGAGC
ACTTTAACCC TCAGTGGCAA GGTTTGTGGG AACTTGAGTT CGGACCACTG GGATTGGCAA ATTCCCCTCT GGGCTAGGGT
TGCTTTAAAT GCTCCCTTCA CGTGTGGGCA ATCAGCTGAG TTTTGGTCCAG TTTTCCTTTC TGCT

SEO ID NO:1397: (Length of Sequence = 370 Nucleotides)

TTGAGTTINI TCAGTGGCAT CCCCTGCTCC CCTGAGCACA CACAGTGTT TCTATTTATG ACTGTAGTGC CAAGCAGAAT
TTCCATGINC TTGCTAGCTG CCCATTCTCA CCCCTCAGGG TCTCATACTT CTCCCTGGAA GCCTCCCAAG CAGTCAATGT
GACAGGGACC AAGTATGTAC AAGGCAACAT ATTGGGTTCA AGTGCAAACT AAGGGAACCA GGGCCTGTTT TTCTAGTTTG
GAAGTTTTTC TTTATCCTAA GAAAAGAGAC AGACCAAAAC CAAGAAGAT AACAATAACT CTTCTCTTTG TCATCACGGT
GATGACATCA AGGTACTGAT ATTAACCAGA AGTTACAACA AGAAGGAATT

SEO ID NO:1398: (Length of Sequence = 307 Nucleotides)

ATCAGCATTA GGITTICACC AAAGTGATAC AAGTCTGAAG GTCTTCATCA GCAGTCTCCC TCATAGTCAG CGCCATACCG
AAGAGGCCTG TCCCTCTCAT AGGGCCTTCC AGCCACINCT TCCCCCACAGG CCTGATTCTN CTGTGGCTGG GAGTGTGGAC
TGATTTGTTA TGATGTGAGA GATCCCNNGG GGTGTGAGCT ACCGCACCTG GCTGAACTTT CAAGGAGAAG TTTGTGCATC
ANITTTCAAA AAATTATGAT ATCAAAAGAT AGCTGTGCCC TACATTTGGG AAAGATACAA AAACTTG

SEO ID NO:1399: (Length of Sequence = 380 Nucleotides)

CTGAATTATT GAGGATGAAT TGATAAAGAC AGGTGTAATG AACTGAGGCC GGGCATTAGA CTGAGCAGCT GACTGTCCCT CAGAAACCAT AACCTTGCTA CCCGCATTGG GCATTGTGAC AACTGTTGAC ATCAATGCAG ACTGCAAGIN AGTTGGCAAA GCTGCTGATG TGTTAGCTGA AGTTGTGATG GGATTGGAAG TGACAAATAC AGTTATTTGA TTTGGGGGCA AGGGAGINGA AATGGAGGAA GAGCTAACAG GTCTTGACAT TACTGGAGGG ATGCTTGGTG CAACGTTAGA ACTGACCTCA CTCAATTCGG GGATGCACAA GGGATGAACA CAGCTCATTT CCTGTNAGGT AAGTTTAGGG AATTAGAAGG

- SEO ID NO:1400: (Length of Sequence = 232 Nucleotides)
- * ATTATAGATA CACACCACCA CACCOGGCTC CTCACATTAA AGTGGGNTTA TGACCATGAA CACTTCGTAT TAATAAATGT

 CTCAGCACAC CCAAGCCTGA AAATCTGATC TAAACCTCCT TAACTTGAAT TCCATCCACA ATCCACAACT TNCTCGGNAA

 AAATNINTCC CAGCTTCTCC TTCCTCTAGC CCAAGAAACA GCCTTAACAG CGNGCGATTT CATTCCTACA CT

SEO ID NC-1401 (Length of Sequence = 349 Micleotides)

AAGCTAAATT TATAATGAAC AGATTGAAGA AAAATAAAGA GCTACAGAAA GTTCAGGATA TCAAAGAAGT CAAGCAAAAC ATCCATCTTA TCCGAGCCCC TCTTGCAGGC AAAGGGAAAC AGTTGGAAGA GAAAATGGTA CAGCAGTTAC AAGAGGATGT

GGACATGGAA GATGCTCCTT AAAAATCTCT GTAACCATTT CTTTTATGTA CATTTGAAAA TGCCCNTTGG NTACTTGGAA
CTGCTAAATT ATTTTATTTT TTACATAAGG TCACTTAAAT GTAAAGGCGGT TAAAAAGACAT CTTINCINGC ATTGCCATCT
TAATATC AGATATTACG GGATGTTAG

SEO ID NO:1402: (Length of Sequence = 338 Nucleotides)

GTAATTGCTA TITGATGTTA TITTAAGAAA TTAACCCITA AAACTITAAT TCCITAAAAC AATCTCAAAC AGAAGAAGCA AAAGCTTGTN CTGTGCTCCA GGAAATAAGA TTCAGCACCA ATGAAAATAA ATTATAGAAA ATCAGAAGAT GGGTCAATAT GAGGGAAAA AACCTAACAT TITAATTGTT TTINCTCTCA ATAATTGTGT TGAACCATCC AAAAAAGTAT GATACAAAAA TAGCACTATA CTAAGAGCCA GATGACATGT CCTTAAAGCC TTAGCTCTGC AAATTATTGG TTGTGTAACA CTAGGGAACA ACACTTAGNC TCTCCTAG

SEO ID NO:1403: (Length of Sequence = 381 Nucleotides)

GGAGICTCAC TITIGITGCCC AGGCTAGAGI GOGANGGCGI GATCITNGCI CACCACAACC TCCATCTCCI GGGITCAAGC GATTCTCCIG CCTCAGCCIC CTGAGCAGGI GGGGITACAG GIGCCCGCCA CCGCACCCAG CCAACITINI GITCTCAGCA GAGACAGGGC TTCGCCATGI TGGICAGGCI GGTCTCGAAC TGACCTCAAG TGATTTGCCC ACCITGGCCA CCCAAAGTGC TGGGATTATA GGCGIGAGCA CTINCACCIG GCCTCTAAGC TTAATCATTI CTAGGCTTTT NATITAAAGI GAGAAACATG TGACTCITIC CTITCATTIG GGACACTITA AAAGGGGTTA TTAAATTGAC CCTAATTACA A

SEO ID NO:1404: (Length of Sequence = 325 Nucleotides)

AGCTCATCAG CTATCATTIG TGITAGIGIA TIINATGIAI GGCCCAAGAC AATICINCIT TITCCAGIGI GGCCCAGGGA AGCCAAAAGA TIGGATACCC CIGACAGGAT TCCAGGATTC TITIGIAAT: NCICAGAGGC CCICIGIGCA TACTCCGIAA GGACTATCCA CATTCTTAT TACITTCATI GGCAATAGGI ATAAAATITT ATITGITGGN TATTTACIG NAATGTTACT TGITTTIGCT TATTTACIGA TIGGGIGGGA GGAAGGCTAA GGATGAATAA ATCTAACCNI TITTAAAAAG GAAAGGCTAA AAATA

SEO ID NO:1405: (Length of Sequence = 349 Nucleotides)

GGATTATGAC TGAACGTCCT CAGCATGTTG GCCTTCACCC CTGGCGGTGG CTCGAACACA AAGATGCGGC CCGCACGGAG CAGATTCACA GGCACCTTGG GGTTGATCTC CATGGTTAGG AAGAGTCGGA AGCAGGCATG CGGCTGCAGG GAATGCAACT TCTTCTCCAG CTGCATCAGC CACCCTGGGG CCAGATGCAC ATTCTTCAGC ATCACCCACC TGCCCGANIT TACAAGCGGT GTTTTATTGC CTTATCTGCT TNGTTAAAGC CTTCTTCAGA GCCGATTGCA ATTGAAGGGA TCTTCGGGGT TCTNCTCGGC TNCAAAAGGTC CTCGACAATG TTCCCCTTG

SEO ID NO:1406: (Length of Sequence = 392 Nucleotides)

GGACTGCCCG TITGITTATG AGACAGGGTC TCATTCTGTC ACTCAGGCTG GAGTGCAGTG TCATGATCAT GGCTCACTGC
AGCCTCGACC TCTCAGGCTC AAGTGATCCT TGCATCTCAA CCTCACGAGT AGCTACGACT ACAGGTATGC CCCACTATGC
CTGGATAATT GINCCITTIT TITTITTTGGT AGAAACAGGG TCTCATTCTG TTGCCCAGGC TAGTCTCAAA CTGCTGGACT
CAAGTGATCC TTCCAACTCG GCCTCCCAAA GTGCTGGGAT TACAGATGTG AGTCACAATG NCCAGCATGG ATTGTCCTTT
TCAGACCCAG ACCAAAGAAC AGGACTTATT TGTCCCAAGA CCAATCTAGG NAAAGTATAA GCTGTGTTGT CA

SEQ ID NO:1407: (Length of Sequence = 362 Nucleotides)

GITAATTGGG NTTCACAAGC AATAATTTCT CCACAACAAA AACCACAACT TGAAGNGAGT TGAAAAGNGN TCAATAGTGG AAACAGTCGC CTCAGTACTT TINCTTTCTG GNITTCATCT CTAGAAATTT NAAGTGTITN AGNCAGAGTC CACCCTTTGT

335

GCAAGGCENG AACCNATGAA TGGACTCCTT GTGTGAATTA TTGCATCTTC TTCCAAAGCA GGTTCATCAA GACTTTCACA GAGATTCATT TTINITGAGA AGTAAGGGTT AATAGGAGGA TAGAATTTGG TTCCNAATCT AGTGNTAAAA GTGTCCAAGC AAATCAAAAA GTAAGATATT TTAGGGGCCA TACCCACATC TT

SEO ID NO:1408: (Length of Sequence = 388 Nucleotides)

CCCCGCAGCA CCACGAGCTG ACCTCGCTCT TOGAGTGTCC GGTCTGCTTT GACTATGTCC TGCCTCCTAT TCTGCAGTGC CACGAGGGC ACCTGGTGTG TAACCAATGC CGCCAGAAGT TGAGCTGCTG CCCGACGTGC AGGGGGGCCC TGACGCCCAG CCCTGCACCA TACGGAGAAA CCAGAACATG AAGACATATG TMAATACCGT CCCTACTCCT GNCCATGTCC TGGTGCTTTC CTGCAAGTGG CAGGGGGGCC TGGTCCCTTAC TAGGGAGAAA CCAGAACATG AAGACATATG TMAATACCGT CCCTACTCCT GNCCATGTCC TGGTGCTTTC CTGCAAGTGG CAGGGGGGCCC TGGGGAGACAT TGATTGTCCC ATMINAATGG AACGGCCCAC AAAGAGCA

SEO ID NO:1409: (Length of Sequence = 348 Nucleotides)

CAATGAACTC CTTAAGCTIT GTTAATATGA GAATGTCTTT ATCTCTTCTT TATTTCCAAA GGACAGCTTT GCTGGTTAAA
ATATTCTTGG TTAAGTTTTG TTTTTAGTAC TTAGCCATATA TCATTCCACT CTCTCCTGGC CTGTAAAGCC TCTGCTGAAA
GATCCACTTC TAGCCTTATT GAAACTCCCT TCTATGTTAT TCCNTTCINC CTCTTGCTGC TTCCAACATC CTGTCTTTGT
CCATAATTTG TAACAGATTG AATATAATAT GAATTAGNCC TCTTTAGACT GAATCTCATT GGAGNCTTTT CACCCTTCTT
GTTTTTGGGT ATTTAINTCT TTTCACAG

SEO ID NO:1410: (Length of Sequence = 370 Nucleotides)

GACTATTIAT TCTGCCTTAA ATCAATGGCA AATAAGTCAA GATGACATIT TGTGAATGTA GACTATGGAT ACACTCCTAA
TAGATTGATG TAGTCATAAA AGGGGGTCAA GTAGATGTT TNCTGTTATG TAAGCAATAA TTTTCCCGTG TCTTATTGAG
TATGGCTAGC GATTATTTAT TACATGCTAG ATGGGTTCTT TGCATGTGGG TTCCATATAG GTGCAGAAAT TTCCTCAGCC
ACTGGAGGGA TTTCGACCAT ATTTGTCATT TGGATGAGCT GTTATTAGAT TGAAATCTAC ACATCATTTC ATTAAAAATT
GTGCCCTAGA AAACGCAAAG CTNTTGCACA ATGGCGATTA AAATTATGGG

SEO ID NO:1411: (Length of Sequence = 385 Nucleotides)

SEO ID NO:1412: (Length of Sequence = 337 Nucleotides)

CCATTCAGAT TCCTCCTGGG CCTCCTCGCC CCATTTGCGA CAGATTTGCT ACCTGCTCCA GCTCAGCGAC CCTTCCCTCT

ATGATGAAGT GCATTGAAGA GAACAATGGT GTGGACAAGA GGATCAGCAG GTTTATTCTC CCCATCGGGG CCACCGTGAA

CATGGACGGA GCAGCCATCT TCCAGTGTGT GGCCGCGGTG TTCATTGCGC AACTCAACAA CGTAGAGCTC AACGCAGGAC

AGATTTTCAC CATTCTAGTG ACTGCCACAG CGTCCAGTGT TGGAGCAGCA GGCGTNCCAN CINGAGGGGT CCTCANCATT

GCCATTATCC TGGGAGG

SEO ID NO:1413: (Length of Sequence = 367 Mucleotidae)

ATAAGTGGAG TGAAGAAATT AATGCATAGT TCAAGCCTAA ACAATACAAG CATCTCACGC TTTGGAGTCA ACACTGAAAA TGAAGATCAC CTGGCCAAGG AGCTGGAAGA CCTGAACAAA TGGGGTCTTA ACATCTTTAA TGTGGCTGGA TATTCTCACA ATAGACCCCT AACATGCATC ATGTATGCTA TATTCCAGGA AAGAGACCTC CTAAAGACAT TCAGAATCTC ATCTGACACA
TTTATAACCT ACATGATGAC TTTAGAAGGC CATTACCATT CTGACTTGGC ATATCACAAC AGCCTGCACG CTGCTNATGT
AGNCCAGTCG ACCCATGTTC TCCTTTCTAC ANCAGCATTA GACGGTG

SEO ID NO:1414: (Length of Sequence = 360 Nucleotides)

GTATACAGCG TGGTCCAGCC ACCCGACAGC GAGATGGGCA TITTAAGAAA CGCTCTCGGC CAGATCTCCG AACCAGAGCC AGAAGAAACT TNTACAAAAA ACAGGAGTCA GAACAAGCAG GGGTTGCTAA GGATGCAAAA TCTGTGGCCT CAGATGTTCC CCTCTACAAG GATGGGGAGG CTAAGACTGA CCCAGCAGGG CTGAGCAGTC CCCATCTNCC AGGNACATCC TCTGCAGCAC CCGACCTGGA GGGTCCCGAA TTTCCAGTTG AGTCTNTGGC TTCTCGGATC CAGGCTNAGC CAGACAACTT GGGACGTGCC TCTGCATCTT CAGACAGAAT TNCTAGCCTG CCTNAGGAAA

SEO ID NO:1415: (Length of Sequence = 314 Nucleotides)

CTCAAACACA GCATTIGAAG TCTTAATATT TTAGTACATA CTATACTATC TCINCTTACA ATTGTTTTTT GITAAAGAAA CCATGTTTTT NATTCTAAAG AGITTCCTTT ACTGTGGATT TTACTGATTG CATCTTTGTT GATGGGTTAA GATTGTCCNN ATTAGACAT TAGTNCTTTC AATGTGCTGT ATTCAGTGCT GCCTCTGGGC TCCTAAACTG TGGAGGGCTG TTTGTCCCTA TTACTAAATGG GGACAGATTG TCCTGCTTTT TAATTTTCAA TGCCTGACTT TTACCCNCTA ACTTTTCCGT AGAT

SEO ID NO:1416: (Length of Sequence = 370 Nucleotides)

TTCCATTITT GCICCITCIC AGGATAATAG CAGACCGGIG ATCACAACIT TAGTITIGAT GAGATAACCI CCITATCIII TAAAAATGGI CICIATTATT TTCCAAGAGA AGACCAGTAA ACACTAAACA CCIGCCITGA TCICAGIGIC TTAGATGITT TCCIGITTCT CCITIATCCT AGCAAACICC CCAGGITGCI ATTCITATTC CCATTITATA GATGGGCAAC TGGGTAAGAG AGGTAAGCTT GGTGAGGICA CTGAGATAGI GGGGAAAGGA GCITGGITCA CATCAGGIAT GCATTCCCC AAGGITCCAC TGGGGCATCT GAAGGAAGGG GTITCIGGAA GTGCAAAATA TAGGGTACTG

SEO ID NO:1417: (Length of Sequence = 365 Mucleotides)

GACTOCTICG CCAAGGGAGC CATCAGCACC AGITGITCCA GAGCAGCCAC AACACCAGIG ACCICCCTIC TGCITCCGGC CAATCCCGGAC AGAGCCICTI CCCGAGICII GAGCTCCIGG ATAGCTGCCI CAATAAAGCA GGACTCGGGA GIGIGCITCT CCTCTGCCAG CTGCTGCTCT AGIGCTACTI TCTCCTCCAG AACTACCCGG TGCAGCACCT GCTCCTTAGA GGCCAGCAGC AACTTGGAGT ACTGGCTGIG CTGTTCATCT CCTAGATGAA TGGGATGGIC TACATTCATC CATTTGGGCAAA AGCCCACCAAC AACCCCTTTT TTTCCCTCTT CAATCAAGCT GCAAT

SEO ID NO:1418: (Length of Sequence = 354 Nucleotides)

CCAAATCCTT AAGITTACAA AGCTGTTGGA AAACTTTGTG TCCTGATTTC AACAATCACG CTTTGTTTGA AAGATGAGCC AAGCTCACAG ACACTAAATT TTATGTCATG CCATAAGCTG GAGAGGAGCC ATTTGGCTAC AGCTGCGGAA CTTCATTGAG GAGCAAATGA AAGGCACATG GACGAGCACG CTGGTGCAGT TCATGTTCTT CCTGCCTGTG AATTGAATAC TGTCCTGGTA GCAGTTTTGG GTCGGTCAGG AGCTCAAGGC TGGTTTGTGT GGCTGACTAC GGATGAGCAC TGAAGTTGCC TCAAAGAATT AAGGGTGTC CACANCAGCC TCTTGGGGTC TTTT

SEO ID NO:1419: (Length of Sequence = 363 Nucleotides)

GTGGAAAACG TCGAAATGAT GTGGCCACTG AAAGAAATTC AAGACAACTG AAACAACTGC AGATTTCCAT TTTCACCTCC TGTTTTCTTA TGAACAATAA CATTGCAGAA GGGGAAATAT CAGAAAGTTG ATTGATTTTT AACCCAAAAA TAGAACTTTT TGTAAGCTAG GAAAGCATCT AAAATTAACA AGAATACAAA AATGCACTTT TGTTTACATT TGCTCTATTT AGATCTTACA AGAGATTATG TCTTGAATCT ATCCTGACTT CAGCAAAAGA CAAAAGAACG TTGAAAACAT CCTATTTCCA AATCGTTTAC AAGGAAGTTAC CTAAGGAGNC TGACAGATTC AACGGCTGCT ACC

SEO ID NO:1420: (Length of Sequence = 326 Nucleotides)

GAAGATTTIC TAGAAGCAAA TAGIGCCACC ATCCGTCATG AGENICIGIT TCTATAACGC TIGINIGICI TINAGACTAC
GIAGGIGGIA GCTIATGAGI AGIAATGINC TITIGITAGI AAATGICACC AAATAAGCAA ATAAGAGAAA CATGAAGGCC
AAAAACTGIN TIACIATICA GGAGAAAATG GACGGITTAG CAACAATACA ATGIAGACIT CAAAATATGA AAAATCAAGG
AAATINCIGI CATTGICTIT AAGGGCCTCC AGAGAAGIAT TAATTIGTCC TITIATGIGAA TITIAATGAGA TCATGIGAAA
TGIATG

SEO ID NO:1421: (Length of Sequence = 294 Nucleotides)

ACCCAGTACA GGTACTCTCA CAGGAGGCAC TCAGCAGGGA TGTAGTGACA GCAGGTAAGA NICCACCTCT NICCCTGCCT GCNCCTGGGA TCCAGTATTG GCCCATGTAT CINCCCCATT TCCTCAGGCT TCCTGGACTT TINITGGAGG GAAAGAGGAA CAGAAAGAGG AGCAGGCAGG AGAAGCAAGA GCTCCCGGGG GCTATGAAAG GTAACATACC TGGAGAGTTT NGGGAGACGG CGGCTTGTNA GAGACAAGGG GAAGAGACAG AAACAGGAGT ATTCTAAGAA GCAT

SEO ID NO:1422: (Length of Sequence = 306 Nucleotides)

GAAGGGCATA TITAATAGCT GCTGCAAACA TATGGAATAG TGCTTTAATC AGTGGTGAAC AAGAAATTGC CTGTTGTTGG TTATAAAAAC AAGGGACATT AATGINCTTG TTCTTGTACC ATAGTAATGT GNAAAAAAAA ATAGTGGTTG NAATGGTGT TAATTTGTAC AGTTTGTGC AAAGTAGAAT GGGNCAGATA TTTTGGTGGA TAGGCTTTTG TCTTAGTTAT AAAAATTAGG NCATTTGGTA TGATAAAGGC NGAGAATCTT AACAATTGGG CACTGGCCCA GAAAATTNCA GGGTGC

SEO ID NO:1423: (Length of Sequence = 274 Nucleotides)

TETETETETE TETETEAGAA ATGGGGAAAG ACTGGTCTAG ATAATATTTC AGGTACCTTC CAACACTAAA ATGGTATGAT TCCCAGCTTA CAAAAAGCAA ACTATTTTAA TATTCACCAC TCAATATAGT GTATCAAGCT CTCGGTTTAT GTTTAAGGGC TTAGGGAACA GCAGCAACTA TTCGTGGGCA ATTAATNCAA AAACTCATGT TACCAAAAAG GCATGTTTAG GACCTGCAGG ATAGTGAAAA AGCAAGAACA GTCT

SEO ID NO:1424: (Length of Sequence = 297 Nucleotides)

GEAGGATTAC TIGAGCCCAG AAAAAAAAA AAGCCICAGG GGITTCGGIG AATGITGIGI GGACITCCGI GAGAACAGAC GITTGAIGIG AACIGANITC AAGGCIGATA CAGCCCAGAA CCAGGNACAA GGIGAGAAAAC TGCTCGITTC CGGGAGGCAG GACTICCCIAA CCGGGAGGCA CTGCAGINCA CITTCTGAAA CAGGITTGGA GGATAGGGAA ATTCCTGNCA GCCCGGGGGG ATCCACITAG TITCTTAGNA GCGGCCGCCA CCGCGGIGGA AGGCTCCAGC TTITGTT

SEO ID NO:1425: (Length of Sequence = 276 Nucleotides)

ATTITICAA GGATGGAAAG GICAGAGAAA AATAAAATAA AACATCITIC AATAGTCITI CCTGGIAAAA GCAGCGICTC TMTGGGCIGG GGAGTAAAGG GIGIGGGGCA AGGGGAGTGG GGAGAGGCTG TAAACCTTCC CCCAAACCCC AGTITTAGAT CCTTTGGTTT CCTTCTCCCA GAAGATGGNC AGAAGGGCAT NGTGGGNAAC AGCAGGGNGG AAAATATGGT GATGACAAAC CCCAGATGAT CAAGGGGCTG ATGCTCCTGG GGCCCA

SEO ID NO:1426: (Length of Sequence = 295 Nucleotides)

....

TAGTGGCATA TGGACCGGAA AGGGTTAATT TAAAGGGGGG GAACCTCAAA AGTTTTTTTA AAAAAGAAAC TTGTCTGCCA CAGTATGTTA CCAGTGTTAA CCCTTCTGCC AGTTAGCAAA CTTTTGCCTT AAGCCTTTTT CCTCTAGGAT ACTCCCCATG TTTCGGTAAT CTTGGGCATA CATTTTTTAA GNATGGACCT CTTTGCCTTG TTTTGTTTTC ATGCTGCTGT ATGTCCAAGT ATTGTTAATT TCATAATAAG ACAAGAGTTG CTTTCTTTTT TAATTCCTTT TTCCC

SEO ID NO:1427: (Length of Sequence = 207 Nucleotides)

TCAGGAATGA TAGTATCTGG GATGAACTCT TCTTTAATAA GATTCAGGCC AGINTIGGIG GGIGINIGCG GATGATTGTT ACTGGGAGGCAG CCCCAGCATC ACCAACAGIT CTGGGAATTT CTCCGGGCAG CTCTAGGGIG CCAGGITTAT GAAAGGITAT GGCCAAACTG AGIGCCACAG CTGGATGTAA CCTINCACCA CTCCTGG

SEO ID NO:1428: (Length of Sequence = 223 Nucleotides)

TAACATTCTC TCCAACCTCC CCAGGTCCCA TCAGTGTTGA GAAGGAATCT AGGCCAGCTC CTGGGAGATG CCAGTTAAGC CGCTTTGAAT CCTGTGCCTT TCCAATTGNC CCTTATAGCA GTCGATGTCA GGGATTGGGA CAACTTTCAA AACAAGTCCA TCAAAGTCCC CATGGGCACT AGGGGCTCTG GGAACCCAGT GTCGAGAGGC TTAGAGNCAT TGC

SEO ID NO:1429: (Length of Sequence = 222 Nucleotides)

AAAACCAAGG AGCAAAGGGG AGACAGAGAG AAAAGTGGGA TGGATTCAAA GACATTGCAA CATAGAACTN ACCGAACTGG CTTGTNTGAG GTAAGGGNGG CAGGATGACT CACAGGTTTC TGGGATTATG TGCAACAGGT GGAAGGTGAT GCCATTAGCC AGAATAAGGC TGTAGGCTNA AGGGGAGTNA AACTGGTTCT GGGGGTATAA CATTGATAGG CC

SEO ID NO:1430: (Length o: Sequence = 246 Nucleotides)

CAAAATTICC TGIATCCITT CATGGGTIN CITTIGITIG TITIGGIAAG AACATTIAAC ATGAGATGIA TCITINAGIT GITGITGIGG TIGANCITIT TIAGATACAT AGICTCACTC TGITACCCAG GACTGGAGTG CCAGTGGACA TGATCCACAG CTCLCTGACA GGCCTCAAAC TCCTGGGACC CAAATGAATC CCTCCCACCT NCAGCCCTCC CAAGTAGGCT AGGGACTACA GATGIG

SEO ID NO:1431: (Length of Sequence = 364 Nucleotides)

CTINCCACTO GATGATGOTT CTATAATTIT GCCCTITAAC AGAAACTITO AAAAGGGAAG AGITITTGIG AATGGGGGAG
AGGGTGAAGG AGGTCAGGCO CCACTCCTTC CTGCATTGIT TACAGTCATT GGGAATAAGG CATGGCTCAA ATCGGCCACA
GGGNCGGTGA CCTTGTGCCC CAGGGTTTTG CCCCCAAGTG CCTCCATTTA AAAGCATTAA GGCCGGTACG GCATCTTCAA
AACAGAGGGC TGGCATTCGA GGAAACCCTT GCTGCTTTAG TCCCGATAGG GTATTTGAAC CCCGCNTATA TTTTAAGGCA
TTTTAAATTC TCTTCCCCCC ATTTTATTGA CTTTGAACAA TTAA

SEO ID NO:1432: (Length of Sequence = 208 Nucleotides)

GIGAGINAAC ATGGATGGAA ACAAATTATT AGGITGINCA AAGTGAAAAA CACCAAAAAT AAGATTTAAA AAGAATGTCA GGTATCCATA GAAAAATATT AATAGGICTA ATACATATGI AAAANTIGGC GICCCAGGGG QNAGAGACTG NAAAGTTATA TITTINNATGG CTGAAATCCC CCCAANTITA ACATAAAGCA CAACATTT

SEO ID NO:1433: (Length of Sequence = 274 Nucleotides)

GGAAGGITTT TAATGCATGA AGTATACTIG TGATCCIGGA GGITGGAAAA GATTCAGTAA AGATAAAGIT TGGCAAAAAT GATTCINICC CIAGGATTIG GGGATATGIA AATCAAACCA AAGGCACATT CIGCAGCTCA CAGCAACCIT CATITITITGI CCTAGATTGA GITATCTATC AAGAATCATT CATTCCCTCT CAGCCCTTGC AACTGTTTCC NATGACTTTG GACTTGGCCA TGCAACTTGC TITGGCCAAT ACAATGTGAG TTAA

SEO ID NO:1434: (Length of Sequence = 249 Nucleotides)

SEQ ID NO:1435: (Length of Sequence = 201 Nucleotides)

GAATGGGCC AATGGCACTC ACTGINTCTT CAGGCCCCCA CGGACGGCAT GCCTGGGGAA GCCTAGTCTA CTTACCATCA GCACGTTGAT CINTCACACA GCATGGAGCC ATAGTTTACA AAGGACCACG GCAGGTCAAG GACAGGCCAC TAAAACTTTT GGTGCTGGGC ACATNACCCA CCCTCACCAN CATCAAAGAC A

SEO ID NO:1436: (Length of Sequence = 312 Nucleotides)

SEO ID NO:1437: (Length of Sequence = 294 Nucleotides)

ATTCCAATGG TAATACTAAC GAATTGTGCT ATCTAAATAT TGGATAGTAA AAACGTCAAC ATTTAGAAAA TGTATATCAC
ACAGGGAACC AATATTTTINC AAATTATCCA CATCTAATAT TAGGCAACCA CGCGCAANAA AAGACACGTT CAAAGTACAG
GAGAAATGGA TGGATTTTAA TGTGAGATAG TACAAGANGT TTATTGATAT AGTTTCAAGA TTCCATATTG TAATAAACCT
TTAANGAAAC TTTCACTTCT TGAGTTTTGG GTATAGGAAT CCAAAAAAAA AAAA

SEO ID NO:1438: (Length of Sequence = 311 Nucleotides)

GGCCCTTIGA CTTTGTGAAT GAGCACAATG AAATGCCGCC TACTGATGCT TCTNATGATC AGAACTCTTT TTTAATAAAA
TAAATAACAT AAATCGTTGA ACATAATGTT CCNGTTGAAT GCAAANCAAA AAAAATATGG NAAACATTTT GNTAAAAATTT
TTTCCNGNTA AAACCATGAA CANTGGCTAT GATGAAGGTT ATTACATATG GAAAAAAAAC TCACACAAGC ATATTTGNAT
TTGGCTTGAA GGGAACCCAT CATTAAATGC AANGCTAGGG ATTCTTTTNG AAGCAGTTGA TCCTCAGGTT T

SEO ID NO:1439: (Length of Sequence = 265 Nucleotides)

OGTGACACAG TTGAAGGAGT OGCTTAAAGA AGTOCAGCTG GAGAGAGATC AATATGCTGA ACAAATAAAA GGAGAGAGGG COCAGTGGCA GCAGAGGATG AGGAAAATNT OGCAGGAGGT TTGCACATTG AAGGAGGAGA AGAAGCATGA TACGCATCGG GTAGAGGAGC TNGAGAGGAG CTTNTCCAGA CTCAAAAACC AGATGGCTNA GCCACTGCCC COGGATGCCC CAGCAGTNTC CTCTGAGGTG GAGCTNCAAG ACCTT

SEO ID NO:1440: (Length of Sequence = 241 Nucleotides)

GUITTACTOT TGTGAAGATA GCACTTTAAT CCTAAATEAG CATGTAAGT GTGACAGATC CTATATCAGT TTTTATAATT GAAGCAGATA GTAATAACTA GATTATTGAC ATTTTTCAGT CATGTGTTCA GCTATTGCTT CAAACTTGCT CAAATTATAC TTGENATTIT ATAGTGITTI ATTTATTATA TACTCINCIT GTAATAANNI GGTAATCTAG TTTCCAGAAT CATGCAAATA G

SEO ID NO:1441: (Length of Sequence = 247 Nucleotides)

GACCCCGATA TICCGGCATC ACATAGATAT CCTCCAGATA AANGGTGCGI CCCTTCCATG TACTGTAGAT GAAATAGTAT ATCCCATAGC CCACCACGCA GGGCCCCAGT AGCTTCCCGG GCGCTGGAAG AATCTCTGCT ACCAAACAGT GATAGAAAGG ATTGINICCA AAGCCATCTG CTCTCAGGGC TTCTTCACTG ATAGGAGTTT TTTTCAAGNA ATAATCCATG CTAAGAATGG GGTATTT

SEQ ID NO:1442: (Length of Sequence = 233 Nucleotides)

GATTACAGCC AAGTTCATGA ATACAAATAA AATAGCAATT TCCCTCATTC TCTCTTTTGT TTTCTGNTCA GAGAAATCAG
GAGATGGGAG CATTATGCTC AGAAACCGAA GAGCTCTTCC AAGAGCTCCA GCTTAGAGTC CAGGCTTCCA GAGCATGCAG
CCTCCTAACA CGTATGTGGT CACATGTGCA AAGACCTNTA TTACAAAATA TTCAGAGCAG NATTTCTNTT AGG

SEO ID NO:1443: (Length of Sequence = 288 Nucleotides)

AATAAACAAT GIGCAGGTT TTATAACIGA TCGGAAGAAG GIIGACCCNC AGITATCACC TTTAAAAAAT GGICTTAGIT
AGGCTTTCIC CCITTGICCT TTTCCAGAAG AAACITGGAG TCIGICAAAT TTCACAAAAT ACCCTGITGA GATTTTCCTT
GGCTTTCATA AGGGIGAATT CACAGATTAA TTCGGAAAAG AATTTACGGC TTTCTAATCA AATTGITCCT TCCAGGGGNT
TTTGTCNTTA TTTAGGNCCT TCTAAAGGIT AACCCTAACT TTGATTAT

SEO ID NO:1444: (Length of Sequence = 208 Nucleotides)

GGAACTGAGT CACAGGGCCA AAGCCCCCTT TNCCTCACGT GAAGCAACTC AGTAAGATGG CGGTGCAGTG AAGCCTATTC CCACACACCT CGGCACTGAT GGAGCAGTCT CCAAAGGAAG GCTGAAAGGA CAGCAGGTGG TTGCCTTNGG GTCCTTCCTT CCCATANCTT TAGAGTGCCA TTTTTCAGCA ATGGGTAATA GCATCAAC

SEO ID NO:1445: (Length of Sequence = 239 Nucleotides)

CCCCGGICIC INGGACACCA TITICIGCCG CIGGACGCAA GGGITTGTGT TIAGAGAATC AGAGGGATCT GCATTAGAAC AGTITIGAAGG TGGCCCCIGT NCIGITATTG CACCIGINCA GGCATTTCTT TIGAAGAAGC TCCTGTTTTC TTCGGAGAAG TCTTCTINGC GGGATTTTTC AGAGGANGAG CAGAAGGNAC TCCTTTGTCA TACCITGTGT GATATTTTAG AAAGTGCTT

SEQ_ID_NO:1446: (Length of Sequence = 243 Nucleotides)

TGCAGGGAAT TINITGATGC AAAACCAGGA AACAATITAT CTCCACTGGG AATACTTTGA AGAAGGGATT AGAGCGGGGC
TAGGGCAGGG AGGATCINTA AAAAACAATA TTTGCCAAAC TAAAAACACA TAGGCACACA TGGGNATTAT TTTACTITCA
ACAAGITCTG AAAGTAGTAA CAAAACCAGG GAGAGITAAA AGAATAATTT AACACTNA'IG NITCAGGAAT GCTAAAGGAG
ACC

SEO ID NO:1447: (Length of Sequence = 371 Nucleotides)

AGITATAAAT GAACATCIGI TGCCTACTTA ATAGGTCATI GAGTAGCTGI GACCCATTCI TAATTIGTAT GTAAGCATAT
TITTIACATA TITGIATCIA CITCATTITC CCTIGAAGCI TGCCAAATTG GTACACTTCA GITTIAACTG ATGICTCITA
TATGCTGTAC CACCITCITA AAAATTGAAT TATCTTTCCT TCCACCTAGA TIGITCTCAA AGCATTTGIT TTTTGCTGGAC
TTTCCACTCT TGACCATAAG ATGGTAGCAT TCCCTAAGGA TATTGCAGCA CAGTCTAATT CCACTGGTTG TCATCTACAG
TTAAATCGCA AATAAAAAAT AATAATAAGC AGCAACTGAT TGCTCAAGTT G

SEO ID NO:1448: (Length of Sequence = 366 Nucleotides)

AATTTIGIGT CCTGTAGGAA ATGCTTCCTT GGGTGTTGT ATTATAGCCC AATCCAAGTC ATCCCTGAGA ACATCCCCAG
GTTGTAAGGA TTAGTCAGAA GTCATGATGA CTGTCCTATA TAAATATTTG GCCTATTAAC TAAAATTAGT ACCTINCCAT
TTCTCCNCTT TCTTGGGCGG GGCAGCGGG GAGTGCAGGG GAGGGGAAAT AGGGAACGIN CAATTGINTT TTAAGTAATG
CTCATAAAAAT TCTTAGNCAA AGATGATCTT GCCCTCCACC TTGGTGACCC ACCGCATACG GGGTACATCT ATCTGGCCTG
TCTCTAGGCC TAGACAGAAG GAACAGGAG GGTTATTGTT AACTTT

SEO ID NO:1449: (Length of Sequence = 234 Nucleotides)

GITGIGGGAG GGACCCGGIG GGAGGIAACT GAATCATAGG AGCAGITTCC CCCATGCAGC TGICGNGATA GINAGITTCT CATGAGGATCT GCIGGITTTA TAAGCITCIA GIGITTCCCC TGCTGGCACT CATTCTCTCT CCTGCCACCC TGTGAAGAGG TGCCTTCTGC CATGATTGTA AGTTTCCTGA GGCTTNCCCA GCCATGCAAA ACTGTGAGTC AATTAAACCT CTTT

SEO ID NO:1450: (Length of Sequence = 220 Nucleotides)

GCTTTINCTC TCCCTGTTTT GTTTTGTAAC CTAAGGAAGC AGAGCCTCTG AGACCACACA CAGCAGCGTC GCCCGTCCCC
AGAGGCACCC CGGCCAGGAC GGGCAGGAGA GGAGACCCCC GTTCCTGCAT GCNCTGTCGC CCCGCCACGG TGNTCTCCGC
AGGGTGAGGC AGGAGGGTGG GTGGAGGCGC CACTGNTCCT CAGCTGGAAG GGCGGGGCCAT

SEO ID NO:1451: (Length of Sequence = 403 Nucleotides)

COGCTGITCA CCTACGGCCT GATTAAACIT GCCTTCCTGT CCTCCAAGAC CAGATGATGA TITATTCTCCA CCGTCTAAGA
GACCAAAGGC CAATGAGCTA CCGCAGCCAC CAGTCCCGGA ACCCGCCAAT GCTGGGAAGC GGAAAGTGAG GGAGTTCAAC
TTCGAGAAAT GGAATGCTCG CATCACTGAT CTACGTAAAC AAGTTGAAGA ATTGITTGAA AGGAAATATG CTCAAGCCAT
AAAAGCCAAA GGTCCGGTGA CGATCCCGTA CCCTCTTTC TAGTCTCATG TTGAAGATCT TTATGTAGAA GGACTTCCTG
AAGGAATTCC TTTTAGAAGG CCATCTACTT ACGGAATTCC TCCACTGGAG AGGATATTAC TTGCAAAGGG AAAGGATTCG
TTT

SEQ ID NO:1452: (Length of Sequence = 353 Nucleotides)

TGCCIAGAGA GGGGCCGGGA TITAGAGAGC TGTTCTTCTG CCTATCTGAT CGCCTCCTCA GACACTGATC TATTAGTCTA
GTGCTGCAAT TACTTGGATT GTAATGTTTC CTTGCAATTT TTGCTTTTCA AATTCTTTTC ACCCTAAACT GTAAATACGC
CAGGAGTAGG TAAAAACTTA CAGGTAAACA TTGCCAAGAN ATAAGGATTT TNATGTCTTC TGCTCAGTGG CATAACTCAA
ATCACATGAG ATAGATTTCT TTGCATCTGT CCATTGTATT TCTCTGAGGC TAATTTACAG CACTTTGTCA CGTTAGGNAT
TTTTTTTTCCC CAGTGCTGCT ACTCTCCAAC TGG

SEO ID NO:1453: (Length of Sequence = 258 Nucleotides)

GITGCCCCTN CTGTCTTTCT GINACCCAGA GAAAGCTTCA CAAGCATGCC TGNAATINAG TIGCACCATT TTATTACAGC
TGAAAGANIT GANTGTAAAG AAGGAAGTTT AATAGANCAT ATAATNCAGC AGATTTATTG ATGGGGAGGT ATCTATTGTA
GITTGGCCAG TGAAGGCAGG TCATAGAGGA AAATTTAGGT AAGTCGGATT TNCTTTAAAA AGAGGCCCAA GAGTTAGTAC
CTCAGGATTT TGTTTTCT

SEO-ID NO:1454: (Length of Sequence - 328 Niclebrides) -- "

GAGATGGAGT CTTGCTCTGT CGCCCAGGCT GGGGTGCAGT GGCGCGATCT CTGCTCACTG CAAGCCCCGC CTCCCAGGTT CACGCCATTC TCCTGCCTCA GCCTCCCGAG TAGCTGGGAC TACAGGCGCC TGGCACCACG NCCAGCTAAT TTTTTGTATT

TITGGTGGAG ACGGGGTTTC ACCGTGTTAG CCAGGATGGT CTCGGATCTCC CGACCTCATG ACCTGCCCGC CTCGGNCTCC CAAATTGCTG GGATTACAGG CGINACAACC GCGCCCGGCC GGTAGCAATA GTTTTAATTA AGGTCTTAAA ATCATACAAA AAGGAATT

SEO ID NO:1455: (Length of Sequence = 342 Nucleotides)

AATTTAGGTA GATTAGCATT CCCATGTAAC TTACCAGAAT CAGAATGAGA ATTCAGAAGT CACCTGANIT GGCCGGGCAT GGGGGCTCAC ACCTGTAATC CCAGCACCTT GGGAGGCCAA GGCAGGCAGA TCATCTGAGG TCAGGAGTTC GAGACCAGCC TGGCCCAACAAT AGTGAAATCC CGCCCCTACT AAAAATACAA AAAATTAGCC AGGCACCCTG TCCACAGGCC CCCACACAGAC TCGAGGGGCC CCCATCTCCT GTTCTGAACC CAACAGGGTG GTCCCACTNT GGGACCACAA ACCAGGTATG ACTGTTTNAG AAGCAGGCTC ACTACCAGGN TA

SEO ID NO:1456: (Length of Sequence = 296 Nucleotides)

ATCITIGACC TATTAGGIGA ACAAATGAAC CICACAGGAC ACACAGTATI TITTAAAGGC AGACICGCIC TCITITITIGC CAGINAGCAG TICIAGGIAA CCAAGITACA CACIGIGGGI ATTCCIGCCI GCCICITGAA TACAAAGGCC TAGITCAAGT GIIGCITTIT TNATTICAAA TCAATTTITI CITCITICCI TITTGAGATA AAACTATTAA AAGIACIACI ATATATATAA AANCTCAAAT CAACITITCG GCCICCTCCI CGIGTACCAG GGAGTATATI CIGACG

SEO ID NO:1457: (Length of Sequence = 314 Nucleotides)

GAGGATICAT AAGTAGAATI TATAAAGAAC TCCAAAGAAT CAATAACAAA AAGACTGGCT ATGCCTITCG NAGAGCAGCT GCTGTCCTGG AAATCAGAGG ACAGTGAAGG GAAGTCCGAA GATGAGCCTG ACACCATTCC GACATCCGTC CTCCTGCAGG TCGTGGGAGACT GCTAGGAAAC TTCTTNTGGA CCACGGACAT GGCAGCCTGC NTGAAGGAGC TTGTTTTCCA TCTCCTGGCA GAGCTCCTAC GCACGGTGCA CACCCTGGAG CAGAGGCCGC ACCCCGCTGG CCTGTNCTCC TCANTCGCCC TCCA

SEO ID NO:1458: (Length of Sequence = 254 Nucleotides)

GITCCAGTCA CAGATGITIC ATTATCACTA TICAATATTA TIAAGCATCI AATAAGTATA AGGATGCATG AGICAAGGGT CCCTACCTTC AGGICGGAAG CAGGAAGAG ACCAGATCCI AGAACAATAG GACATGGTAC CCGCTGCCTA GACGAATTT AGAATCCGGC TGGGGTGAAG AGATTAATGA GCGAGTCATG CCATCAATGT GCTGTAACTG AGGTCCTAAA AACCACCCAG CCGCGACACA AACT

SEO ID NO:1459: (Length of Sequence = 343 Nucleotides)

SEO ID NO:1460: (Length of Sequence = 348 Nucleotides)

ATTGICAACA GIGITITIAT TIATACCIAC AAAAAGAAAA CAAGATGATG GIATCAAAAG GACAATTTAC AAACTAAGAA
TAGIAACATA GCITTCAGCA TCCTGTGCCI GANCATCACA CATCTACAAG TCTTTCAAGT CITAATGCAA CAGGAATGIN
TYTGGAGACC NGCAAGAACA TCAATAGAGA GCACTGATCC CAAGGAAAAG CCACTAACCT TITTAGATCAG AAGTCCNGAC
AACGNATTGI TAGGGAGGAT TIGGGAGAAG CAGCCCCTTT GCTTAATACA TINGGACCCC TITCCCCTAA GITGAGGTTC
AACCCITGAA TGCAATAACT TGGCATAA

SEO ID NO:1461: (Length of Sequence = 343 Nucleotides)

TGGGAAGATC AGGICITACT TGTTTTCTG TCCCCTCCAG CGCTAGATCA ACACAGIGIT AAATTAGITG AATTTCAGIG
GAGGAGATAA GACAGAAATG AAATCTGTGA AGATTCAGAC TTTCCCCAAGT TAAAACCAGI CTTGAGITAC AGATCAAGAT
GATGCCAGAA ATAACATCAC ACTGAAACAT CAGTCAAATG TAGTCATCAT GGCAAAGGCC AAATGTCCCT TTCTTTTTTT
GCCTCCGCCT GCCTGGGAAT TTAGCATCCC CTAAAGCCAC TCATCTGGGA CAGGATTTTA GGGTGTGAC ATGTTTTTCA
ATCTCCACAG GACCCAGCTG TGT

SEQ ID NO:1462: (Length of Sequence = 335 Nucleotides)

GGCATGGAGC AGGCAATGAC TTGTTCATAG TCGCTGCAGT TATGAGCACC AGCTTGAACT TAGGAACTCT TATAAATTTC
TGTTTTCAAC CAAGTATTGA GTGTCTGCTA TGTGTCAGAC ACTGCGCTAG GTGCTGAAAT CTCACTTCTA CTGAGGAAGA
CAGGAACATA AATGGTGATG ATCATTGCAT TAGAAGTGAT GCCACGGGAA TAGTGTGGGG CCTCTCCAGG GGGATCINAA
GGTAGGGAGA CCACACTTCT CCAGTGGTGG AGAGGGCAGA CAGCGTGTAT NGGGTCCTCA AGGTCTNATT GCAAAGGTCA
TGTTTTAGCT GTTCA

SEO ID NO:1463: (Length of Sequence = 382 Nucleotides)

GGACCGCTTT CGGTTCCTCA GGATAAACAC GAGCATGCCC ACCACGGTGA AGGCGGAGGT GACAAACACC AGCAGCAGTC CCGGGACCAA CACCGAGATG GACACCCAGTG GTAGGAGTTG GAGTGCGTCC CGGTCTCCGC CAACCCAGTG CTGTTTTTAC TGTGCGAAGT TAACGTGGGC GAGATCCTAG CGTACAGCTG AGGGCAGATC TCGTCATTGG AGAGGAGCAT GAAATCCTTT CTAAAGAAGT TCACCGGCGT CTCACACTTN AGGTCGCTCA TCAGCACTTC GGAACCCAAG CNTTCTGNCC ACTTGCTTGA AAGGCACAAT TGTNCAGGAG CACTNCCAGG GGTTTCCGTG GAGGGTCTAT CT

SEO ID NO:1464: (Length of Sequence = 187 Nucleotides)

AANGACCICA TITCAAAGAA GAGCCGTCTC CIGACAAGGG ACGITTCCCA GAGAGGAGAC GIGITAGIGC AACAAAGACC AGGCCCIGGN AGCCACGAAA GCCCTCCAGA TGCCTIGAGG ACGCCGTCIN TAGCCGNGIG GGCCACGNCC GGGIGGGGAC AGACAATGAC AAGAGGCAAG ACAGCCG

SEO ID NO:1465: (Length of Sequence = 276 Nucleotides)

TTIACACAAT CAGTATAATA CTGATAGGAA AACTTGACTG AGTTCAGAAA ANGAAAACGA AGTAGAGATC TCACTTGCAT CAGAACAAAA TGTCAATCTA TTAGCAGATA ATATTCATCA GTATTTTTTTG AAAATACAAT ACCACANGAA AGAAACAGTG GACATTTGGA GGGCTTTGAG GCCTGTGGTG GAAAAGGAAT TATCTNCCCG TAAAANCTAG ATAGAAGCAT TCTCAGANAC TTGTTTGTNA TGTGTGCCCT CTACTGACAG AGTTGA

SEO ID NO:1466: (Length of Sequence = 375 Nucleotides)

GEGGITINAC CATGINCCC AGGIGGGCT CAAGIGATCC ACCCTCCTIG GCTTCTCAAA GIGCTGGGAC TACAGIGIG

AGCCACTGIG CCIGGCIGGT TITINITITIT TNAATGAACA TGITGCAAAT CACGCAGAGC ACCININATT CTGCATTINC
TGGGITATAA CAAACATTGI CATCTCTGCC TACATTTAAA AGGCTCTGGI GITATITTAA TATGTCTTTI CAATTTAGTA
ATTAATTCTA ATTITCCTTI GAGCTGAGAT GITATTCATT GITCCCTAG AGTIGCTTTI ATTIGTTCAT ATATGTTTCC
CTTAGCATGI TITTCGTATC TCTTAGTTAT TAGATACCTG AACATTGAC ATTGG

SEO ID NO:1467: (Length of Sequence = 319 Nucleotides)

TGATAAAAGG AAAACGTTTT GATTTATAGT ACCAAGTGCT TAAACACAAG GATAGTGTTA GATTTTCGAG TGACTTTCCT
TTTTGCATTT TTTGGCAGTA AAAGCCAAAC GTTGTATTTG TCCTTTTCAG AGTTGTCCAG CCCTTTTTTC CTTTGTCCAA
AATGATTCTA AATAGAATCT AATAAACCAA TGTAGCATTA TTTTTTTCTA AATGAAGCCC CAAAAAAGAA AAGTGCCTTG
CATCATTTAA AAAAAATAAT TAAATCCTCA TGGCCTCTAA ATTAGGTATG TAGGGCACTG AAAAGTTCTT AACATTTTT

SEO ID NO:1468: (Length of Sequence = 352 Nucleotides)

TITIGGITIAAC ATTICCAAACA TGIATAACCA ATTIAACATG CCTAGGGITT TCITTITATI GGIATICACI TCAGTAACIT GAATCCACAG ATATAAGCAG TATATAACCA GAAAGTTACA AGTIAACACA AATTATACAT GCAAATTICI GITCACAAAG GICACATGIG CAGGIACATG ANITAGAAGC GIGCATCTAG GATIATGGCC AAACTGITTT AAAAATGCAG AAATGTAAAA TTACATCTIG AAAATATGAA GAGATGGICT ACACACTTCA AAAATCAAAT GITGCTTATA CCAGAGATGI ATGTCAATCA CGGGNTTCAA GIGACAAGCA GTAAGGATCC TC

SEO ID NO:1469: (Length of Sequence = 427 Nucleotides)

GAGATGGAGT CTTGCTCTGT NACCCAGGCT AGAGTGCAGT GGCGAGATCT CGGCTTACTG CAACCTCCGC CTCCTGGGTT
CAAGTGATTC CCCTGCCTCA GCCTCCCAAG TAGCTGGGAT TACAGGCGCC TGNCACCGCA CCCAGCTAAT TTTTGTATTT
TNAGTAGAGA CGGGGCTTTA TCATCTTGGC CAGGCTGGCC TCCAACTCCT GACCATGTGA TTCACCTGCC TCCACCTCCC
AAAGTGCTGG AATTACAGGT GTGAGTCACC ACACCCGGCC GGATTCTGTT AGTTTTCTTT AATGCATATT GAGTTTCTTT
AGTTTTAACA ACACCTTAT CTTGGTTGGA CCCAAACTAT TCACTATGTT TCTTGGGGGA NAGCTTNGAA TCTTGGGGTG
GNAGCCAATT AGTAATAGC CAGGGTG

SEQ_ID_NO:1470: (Length of Sequence = 426 Nucleotides)

AGGAGITIGA G-CCATCCIG GGCAACANAG GAAAACCCCG TCTCIACAAA AAGAAAATIT GGITITNATA TITATITGIA
TTAAATITIT TAGAAACATA GCIGGGCATG GIGGCACACG CCIGIAGICC TAGCIACICA GGGGGCTGAG GIGGGAGGAT
TGCITGAGCC CAGGAAGITG AGGCTGCATI AAGIGITGAT CACACCACIG INCIGCAGCC TGGGIGACAG AGIGAGACCC
TGCGACTCCA GACAGGIGCA CACCACCACA CICAGCTAAT TITITGIAGA AATGAGGICT CACTATGITG CCCAGGITGG
TCTIGAACTC CCGGGCTCAA GIGATCCACC TGTCTCAGCC TCTCAAAGIG CTGGGATTAC AGGCATGAGI CACAGIGCCT
GGGCCCAAAT TCATAGICCT AAACAT

SEO ID NO:1471: (Length of Sequence = 372 Nucleotides)

AGAATATTAA AAAAGACCAG ACGCTTAAAG CAAGANITGA AATACCTAGT TGTAAAGATG TGGCACCTGT GGAGAAGACT ATTAAGTTGC TTCCCAGTAG CCATGTTGCA AGACTACAAA TATTCAGTGT AGAAGGACAA AAGGCAATTC AGATCAAACA TCAGGATGAG GTTAATTGGA TAGCGGGTGA TATTATGCAT AANCTTATTT TTCAAATGTA TGATGAAGGA GAAAGAGAAA TCAATATAAC ATCAGCTTTA GCAGAAAAAA TTAAAGTTAA TTGGACTCCT NAGGTTAACA AAGAACACTT GCTACAGGGT CTGCTTCCTG ATGTGCAAGT ACCNACATCT GTAAAAGATA TNCGCTATTT CC

SEO ID NO:1472: (Length of Sequence = 332 Nucleotides)

GGTAGAGACA GGGTCTCACC CTGTTGCTCA GGCTGGTCTC AAACTCCTGG GCTCAAGCNA TCCTTTCACC TTGGCCTTCC
AAAGTNCTAG AACTGGCCAG GGGTGGTGGC TCATGCCTGT AATCCCAGCA CTTTNGGAGG CAGAGGCGGG CAGGGAGTTT
AAGACCAGCC TCGCCAACAC GGTGAACCCA CTCTCCACCA AAANTACAAA ATTTAGCTGG ATGTGGTGGT GGGCGCCTCT
AATCCCAGCC ACTCAGGAGG CTGAGGCAGG AGANTCACTT GANCCCGGGA GGCGGAGGTT GCAATGAGCA GACACGGCCT
GGACGACAGA GT

SEO ID NO:1473: (Length of Sequence = 434 Nucleotides)

GCCITTAATT TGGTTTINCT ATGCCAGIAC AGAAACATCT GGACAACACT CTTGAGCCTG CAGAGGCTCA CGGCCACACCC CACTTCTGCC GCAGGACTGT CTGTTGAGGA GCCGAACCGA TGAGGCACAG TAGCCAGGCC CTCCCGAGGG CTCCAGAAGC TCTAGGTTTA CGGGGTCACC TTCTTGTAGG TGACGTGAAG ATGCTGAGTC ATTGGCTGIN TCGTGGTTGC CATGGAGACC GTCTGCCTCAA GTTTGCCTTC AGAATTCAGC CTGAACTTCC GGGTGATCTG CTCTACGTGG GGCTCCTTGG CGAAGGAGAT CCTGGGGGATG GAGTGGGATG CGATGCACAG NTCCTGCCCG TTCAACTCGC CCTCCTNCAC TTTCCANCAC GGCTGTTTTC TTGGGGTGAC AAAAGGCCAC CTTTTTGGTG TCGG

SEO ID NO:1474: (Length of Sequence = 402 Nucleotides)

GACGITINAGG TGGGAGGITC GITTGAGCAA CATAGTGAGA CCCCGTCTCT ACACAAAAAC AAAAAAAAATA AAAAATTATC
TGAGCATAGT GGAGCATGGC TATGGTCCAA GCTACGTGGG AGGCTGAGGT GGGAGGATTG CITGCNTCCA GGAGTTCAAG
GCTGCAGTAA GCAGTAATGG TGCTACTTCG CTTCAGCCTG GGCGACAGAG CAAGACCCTG TCTCGAAAAA ATAAATAAAG
TAAATAAAGT TGAGAATTIT GTATTTTGGT ACAGAAGGTC TATGCCTTTN AAATGCTCCA TTTGGACACG CTTAGGGCAG
GACGCTCTGA AACTGGGAAG CCTGGGGCCC TGTACANTCT TGGCTGTCCC CTGTACANTC TCCTAACTCT AGAGGGCTGG
TT

SEQ ID NO:1475: (Length of Sequence = 324 Nucleotides)

TIGCATACCT GIGCIGIGIC AGACCAGGCA GAGICATCIC ATTCCACTGG TCTAATGGAT GGCAATTGAA TITTAATTAAC
AAAACTCCTT TGACTTAGIT TCATACTGIG CTGAATGTAA TGGAATCCTC TCTGCCCCCC TTATCTCTCT CTCTTTCACT
CTCTCTCAAC TAAAAATTGI CCTTAACTAA CATCCACTTT AAGAATATTA AAGGCTATAC ATTATACTTA AAAGATACAA
TACAGICATC CCCCTTTCCA TGACTTAAAT TGTATAACAT AAAATAATTA AAAACNTACT TTGGATAGIG ATACACAGTA
TAGG

SEO ID NO:1476: (Length of Sequence = 244 Nucleotides)

GAAAAACCAG AAACTCAAAA TCAGAGTGCC TCTCCTCCTC CAAAGGAACA CAGCTCCTCA CCAGCAACGG NACAAAGCTG
GACAGAGAAT GACTTTGACA AATTGAGAGA GGAAGGCTTC AGAAGATCAA ACTACTCTGA GCTAAAGGAG GAAGTTCGAA
CCNATGGCAA AGAAGTTAAA AACTTTGAAA AAAAATTNGA CGGATNGATA ACTAGNATAA CCGATGCAGA GAAGTCCTTA
AAGG

SEO ID NO:1477: (Length of Sequence = 338 Nucleotides)

ACAACACATA CTIGAAACIG ATTATGACIG INITIGAATG CATTITGATI CCTIAGCIAT GCCICICAGG TGAAAGGACC AATGGCAAGA GGAAGCAGAG GATTCATGCA CTAGAAAATA CTGAGAGAGA TCAGAGTATI CTGTCTACTI CACTGAAGAT ATGGTCTATT GAGGGAAAAC TAATTAACAG TTGATCCAAG GAACAAAAGA ATGCTGTTAT GTGACATTITT GTTGGGAAAC TGACTGTAAT AATAATAAAN CAAATGTCCA GAGGAATGTG TCACATAATT NCAGTGTTTA TGGTTGATAA TTCAAAGGCA TAGATGAATT GGGATTCT

SEO ID NO:1478: (Length of Sequence = 397 Nucleotides)

ACCCITICCE ATTCTGATAA TCTGGCCATG ACTAGCAGAA GCACAGCTAG GCCCAATGGG CAACCCCAGG CCAGCAAAAT

**TCCCAGTTC AAATTGGTCC TGCTGGGAGA ATCTGCAGTG GGAAAGTCAA GCCTGGTAAT ACTTTTTGTC AAAGGGCCAGT

TCCATGAGTA CCAGGAGAGC ACCATTGGAG CGGCCTTCCT CACCCAGTCC GTTTGINTAG ATGACACAAC AGTGAACTTT

GAGATCTGGG ACACAGCTGG GCAGGAGCGA TATCACAGCT TAGCCCCCAT GGTACTACAG GGGTGCCCCAA GCTNCAATCG

TGGGTTTACG ACATTACTAA TCAGGGAAAC CTTTTGCCCG AGCAAGACA TGGGGTGAAG GGACTACAGC GACAGGC

SEO ID NO:1479: (Length of Sequence = 389 Nucleotides)

GCTAGAGNEC CEGCTTGCEG GETTGAETEG CCCGAGCTAA GEGTGCCGAG ACCCAAGGGC GGCGACTACG ACGGCGTTGA
TATCCGTGET AACGACGGCC TCAGCAGGCG GGGAAGATGA AAGGCCCGNT CGAGCTGGGA GATGTGACAC CACACAATAT
TAANCAGTTG AAAAGATTGA NTCAGGTCAT CTTTCCAGTC AGCTACAATG ACAAGTTCTA CAAGGATGTN CTGGAGGTTG
GCGAGCTAGC AAAACTTGCC TATTTCAATG ATATINCTGT AGGTGCAGTA TGCTGTAGGG TGGATCATTC ACAGAATCAG
AAGAGACTTT ACATCATGGA CACTAGGGAT GTNTGGGCAC CTTACCGAAG CTAGGAATAG GGACTAAAT

SEO ID NO:1480: (Length of Sequence = 384 Mucleotides)

CTGAGAGCCA GGAGCTCTTG CGGAGAAGCC ACTGTCTGCA CGCCACCTGC TTCGATGACC CTGCTCTGCC ATCCCTGTGC
TCCAAGGGCC GGGCCCTGCC GTTGCCTGTG CCAGACGGGT CTCAGGGAGA TGCCGGCCAG CAGGTATGCA TGGCGAGGCC
TGGGCATCAA GGCCCGGATT CTATGGCTGC CAGTTTCATT CTCTCGTTGT TTGTCCCCCT AGCAAGACTT ATGAGGTTCC
TTGAGGACAA GACTCCCTCC TGCCACCTGG TCTGTTTCCT GAACATTCAC TGCACTAGCA CGGNCCCGGG ACGCAGNCCT
TGGGAATCAG GCCGTCGGCC ATGGTAGAGC GGCTNGCACT GCTCGGCACC GTGACGGACG TTTG

SEO ID NO:1481: (Length of Sequence = 257 Nucleotides)

ATGICTAGAG CTATTCIGIT TICCCAAGCC ATTIGGTAG TAGGCCCTAA TIGGTCAGTG GGTTCTGACC CCCCAATCCC
TACCTCAGCA GCAGGAAAGG GAAGIGCIGG TCTCCACCTG TNCCCACTAA GGCCCCGIGG TATCCIGGCA GAAGCCTCTG
CATGIATCIN CGCTCTGAGG ATGGGGGITT NAAAACAAAA TAAGACCCTA CGTCCTACTA CCTTGAGCTT GGCTCTAAAA
CCACGGGAAA GGAAGAG

SEO ID NO:1482: (Length of Sequence = 345 Nucleotides)

AATTGAGCTC AGACTAAAGG AATTCITTIT TGACTAAATA GTGATTAAGT TATGATATTC CTGTTGGCCT AAGAACAATG
CCTATGATTT AGTTGTGTA TGTATATTTG TACTTATAAC CAAACAATCG ATTGGGTACA AGTAGCCTTA GGGCAATACT
TCCTTAAAAA CATGTTTCTG ATAAACTAAA GCTTTAGCAT TAACCAGAAG TCATAATTTA ATAGTATTGT AAAAATACCT
CATTTATTTT AAATCCTGTG TTGGGGTAGA GGATTACAGT TGTCATTTCA AATACATGAA TCTCTTGTCA AAAGAGGTAC
TTTGACAGTT TCATGGGAGG TCAGG

SEO ID NO:1483: (Length of Sequence = 344 Nucleotides)

CTGATGTACT GITTTAATAT GCTGAGTACT GITGATTCAA CAACAAACCT TAATGGGTGA TGAGCITTTG CATACCAATA
TGAATTINIC AGCACTTCTG AAAACTGGCC ATCATTITIC AAATTCACAA TITGCTGGAT GTCAGGGAAC AATAGGAAGA
AGAATGAGCG TCAATTTTCA TGTCTTCCTT TGCTTCTTCA CTGGCCTTCC ATAGAAGTAG TCAGAAAAAA ACAAAGCACC
ATCAACCACA CTTCACAAAC AATTCATGTT GGCCTAAGCT TTGCTCAACA TTCATATGAC AGAAGGTAGN ATAATGAAAA
GGGACTGCTG GGCATCACTT TCCC

SEO ID NO:1484: (Length of Sequence = 380 Nucleotides)

TTCCTAAAAG CAGTCTTCC TACAACTTGT ATGCAGTAAG TCACTTAAGC ACTTAAGTGT CATATGGGTA CTTACATGGA ATTAGAGCAC TTCCTGAATG GAATTAGAAA AAGGCAAATT GTGCATACTA CTGATGCATT CATTTCCTAC AGAGATATGA TACCAAGGGC CAATAAGTGA ATAGAAAAAG GGAGGAGGAT TTATTAATGG AATGAGTTCT AACCCTGTCT CTTACCAGCC ATATGACTTT GGGNTAAATA ATCAAACGCC CAATGAGCTC AACTGTCTAT TATTAGGGGA ATTTAAATGA GAGAATGCAC ATTAATTATG CATTGCAGAG TACATGGGAA AATAGTAAAA GCTTAATATT TAATACGGTC

SEO ID NO:1485: (Length of Sequence = 334 Nucleotides)

GAAGGAGCG GGAACCAGIT TCTCACTCTC CTCCCACTTG CIATTGTCAG AAGAGCGAGA TTTCAGGGCA GCAGAGAGCA
TCAGGAGATC AAAAGAAGAC ACTGCTGGT GGTCCCTTAG CAAGITITAG CITCTITINCC TGCTGGGAGA GIATTCCTTG
GGCACAGTGC CAAGTGTCTC TAAGAAACTA GTCATGCCTG ANCITAAGGG CTCGCGGATT CTGGGTGGTG GATTTCCTTA
GGCTTGTCTG AGCCTGCCAG TGCTCTCCTC TGTCGCTCTG ATTTCCATTC ACGCTGAGCA GTCTGCACTN CCTTGGACAG
ACCCACTGGC ATTT

SEO ID NO:1486: (Length of Sequence = 164 Nucleotides)

CTEAAACGGA AAGATGGCGC TGCTGINCTT CTAAGCCTAG GCTTCTTGCA CTAAAGCACC AAGGGCATCG CACACAGGCT TGGCAGAGGG GCCATGGCCA GANTCACCAC CTTCAGACAA GTATGTTGGA GGTCTCGAAT CCCTTGGCAC CCCCAAGCAT GCAG

SEO ID NO:1487: (Length of Sequence = 298 Nucleotides)

TTGAACCCAG GGGCAGAGA TTGCAGTGAG CCGAGATCGT NCTGCTGTAC TCCAGCCTGG GCAACAGAGC GAGACTCCAT
TTCAAAAAACG AGAACCCAGA GGGCTCACTT GCCCCTTCCA CCACACAGTG AGAAGGCACC ATCTATGAGC CAGGAAGCGG
GCCCTCACCT AACAGGATCT NCTGGGCCTT GACCCAGGNC TTTACAACTT CTAGANCCAT GAAAAATTTC TGTTGTTCCT
AGCAGNCCAA ACAGAATTAG AACCATTAAT TTCTATTTCT CCTTTAGCTT AACACTGG

SEO ID NO:1488: (Length of Sequence = 343 Nucleotides)

TIGCTAGTIC AGENTCAATG TCATGGCTGT AACTAATATA GTACATTCGG CAGTIGCAAC GCGAAATGAT CCGCTGGACT
TGCTGGGCTT GCTGTGCCTC ANCTGGCTGG TTCCAATCTG TGGTTGTGGT AACCATGCCG CCCACTGCCT GCCCACTCTC
CATCAGCTCC TGCACAGAGT CCAGACTACG CTGCCGGTGC TCGCTCTTTT GCCCAGGTTG AAGTGCAGTG GCGCAATCTC
AGCTCACTGC AACCTCCGCC TNCCGGGTTC AAGCAATTNT CCCCAACCTCA GCCTTNCGAG TAGCTGGGAT GACAGGCGGC
CGCCACAAACG GCCAACTAAT TIT

SEQ ID NO:1489: (Length of Sequence = 412 Nucleotides)

ATTACCTITT TATAACCCAA GANTGCCATT ATTACACCCG GAACCCTCAC CAAATAAGTA GGAAAACTAC ACTGAGAACA ATTCGGCCCA GCTGTCTCTG GCCCATTTCC CTTTCTACCG CCTCTTGTGC ATTCCAGCAA TCTAACTCGA TGAATGATCT TCCAGTTGGA AAGATGGGGA CTTCACAATG TGCAGACCCA AAGATCTGTC TTCCAAAGGC CAATCACCAC TGTATCCTTC GTTCCTTTAA ATGTCGTTGT TTATTTGAAT ATATTAAGGA ATAATATCAA GGGTAATTAT CTATGTATAA AATGTATGNT TAATTTTTTA GGGGACCATC ATACTGTTTT TCCACAGTGG CTGTACATTT TACAATTCCC ACCAACAATG CACAGGGTTC CATGGTTCCT AT

- SEO ID NO:1490: (Length of Sequence = 356 Nucleotides)
 - ATACCTICIT TCATITAAGC CACCCAGTCT ATGGTACTIC GITATGGCAG CCTTAGCAAA CTAATACGGA TTCCTCATCA
 GGTTCAGATT TINCTAAATA AAATGTGTT GTGAGGGTGG TACAAGCAAC AGTGATATAT TTCTTTAAGT ATTTTCCCCC
 AGCCAAATTC CAACAAGACA ATAATGTCTA ATGCACTGTC TGGTGAATCG GAAAATCTCC TGAATGAAAT AAGAGCCTCT
 AATACCCAAA AGGGAATGAA GTGAGTCATC ACCACAGCCT GTGAATGAAA ATAACTGCTC TGAGGAAAAC ACATGTAAAA
 AATGACACCA TGTGGATTAA ATGGGGNAC ACAAGT

SEO ID NO:1491: (Length of Sequence = 335 Nucleotides)

TTCACTACCA AAACCAGTTA CAACAGTTCC AGCCAAATAA CACAGGCTAC CCCATATGCC ACGACACAGA TCTTGCATTC CAACAGCATA CATGANTTGG CTGTCGGTCT GCCTGATCCN CAATGGAAAA GCTCAATTCA GCAAAAAACA GATCTGNTGG GATTTGGTTA TTCTCTACCT GATCAGAACA AAGGTAACAN TGCCTTACTT TACATTCCTG ACTACCGNTT GGCTGAGGGA TTGTNTAATA GAATGCCACA NAACCAGTCT NAGGATTTTA GCANCCACCA GCTCTNACAA CAGCTCAGGA AGGAGTTGGC AGTNTCTCAG GTGGG

SEO ID NO:1492: (Length of Sequence = 321 Nucleotides)

GACTICATAA AACATCCITT ACTATATITI NAAAGAAAGC AGAAGTAACA GCAATATATG TAAAAGTAAT GNITTAATGN
CTATAAGCAA GNCAAAGCAA TAGAATTGIG CITCITITIGC AGACTGGGGN CAATGAAATG TITAGCTACA ATTINCCCAT
ACAAACATGA AACAATATTC ATATAGNNTA ANCACCCICA CAAATAACTG ATGGGIGATG ANCACACACC AAGTTCGACC
AAAGCAAAAA NTAAACTGAA AATTGITGGG TGGGGITATT CATATTITTAA ATTCAACATG CITGCTCTAT TTAAAAAATAC
C

SEO ID NO:1493: (Length of Sequence = 315 Nucleotides)

GACGGAGCGA GGGGACAGAG CCCAGGGATG GAGGCGGGAT GCGGGGGACA GAGCCCAGGG ATGGAGGCGG GATGCGGGGG AGCAGCTGGT AATGTGCAGA GACTGGGAGA GGGCGGTGTC CAGGTGGAGA GTATTTCAAG GAAGAGAAGG ATTAACAGCG TCCACTGCCG CAGATGGGCC AANCNGAGAT GGGACTGGAA ACCAACCACT GCATTTAGCA TCCTGGGGAC TGCTNATAAC CTTGGTTTGA TGGCTCCTCA AGAAGAGCCA NAACCCTTNA AAGTTAGTTC AAGAGAGAAG GGGNGAAGAG ACACT

SEO ID NO:1494: (Length of Sequence = 405 Nucleotides)

AAAAGTTGAC AAAACATAAA GTATCTCTAG ACAGCAAGGA AATAATTTCA CGAGATTGCT AAATTGATGT CAACACCTGC AGTCTAAAAT TTATACAGTT CAATATGTGT CATTTGATCA CTGGCATGTC AAATATAGAA CAGCTATGAC TTTGCTGGCC AGTAAATTAT CTAGCAGTGA AAATCACTTT TTAGGAGAGT CGCAATCAAA CATTGTTTAA CGTGGGAGCC TATAAAGATG CAAATTCCTG AACAACAGTG TCTAAGAAAA GTACATTGGG TCACTCTGAA CAGGTGGTAT GAACATTTGA TTTAACTGCA AGATCINCNG CINITTACGG GCTTTGTCAC CATCGNATGA ATCTTACATC CGCTGATGAC TNAGAGCAAG CAGGGGCGAG CTGCCC

SEO ID NO:1495: (Length of Sequence = 364 Nucleotides)

CETCIAATGA AGAGCITCGA AACTIGICIT TGICIGGCCA TGIGGGATIT GACAGCCICC CTGACCAGCI GGICAACAAG
TCIACITCIC AAGGATICIG TTICAACATC CITTGIGITG GIGAGACAGG CATIGGCAAA TCCACGITAA TGGACACITT
GITCAACACC AAATITGAAA GIGACCCAGC TACTCACAAT GAACCAGGIG TTCGGITAAA AGCCAGAAGI TATGAGCITC
AGGAAAGCAA TGIACGGCIG AAGITAACCA TTGITGACAC CGIGGGATIT GGAGACCAGN TAAATAAAGA TGACAGCTAT
AAGCCONIAG TAGGNIATAT TGATGCCCAG TTCGAGGNCT ACCT

SEO ID NO: 1496: (Length of Sequence = 370 Nucleotides)

GICTCTTGGA GCAAGGACCC AGITATTCAT CITAATTCTC AGGGGAATCT CIGTAGAGAT GAAAAGCAGG AGAACCAAGG CAGCCTGGTC TCCCTGGGTG ATGAAAAACA GACTAAGAGC AGGGACTTGC CTCCAGCTGA GGAGCTTCCA GAAAAGGAGC ATGGGAAGAT ATCGTGCCAC CTGAGAGAAG ACATTGCCCA GATTCCTACA TGTGCAGAAG CTGGTGAACA GGAGGGCAGG CTACAAAGAA AGCAGAAAAA TNCCACAGAA GGAGGCGGC ACATCINCCA TGAATNTGGA AAGAGTTTIN CTCAAAGCTC AGGCCTTAGT AAACACAGGA GNATNCACAC TGGTGAGAAA CCCTACGGAT

SEO ID NO:1497: (Length of Sequence = 376 Nucleotides)

CACACACATA CAAAATCIGT CCATTIGCCG GAGNAATNIG TATGTATGIN AGTIGGAGGG TATTAAAAAT CAGITTTATT
CCAAAGATIT AAAACTAGAC ATGACTTAAA AACAATTICT GGAGCACTGC TIGCIGACAA TCTCGIAGIT CTCTGCTGCA
TTTGAGTGCA TTTTGIGGCC AGTCCATCAG GGCGTACCAT GGGATTATAT TTGAATGIGT GGIGCATCCT TCCTGGATGA
AGGATGTGTG AGGGACCTTG AACCTCAGCT GTATTAAACT GTAGCGCCTC CAGTCAGTGC ACTAGATGAA ACTTTTAGAC
ANCCTGAATT CTGTTGGGC CNITCTTTTT CCTTTATGTA GGCAGNCTNC AGCATG

SEO ID NO:1498: (Length of Sequence = 281 Nucleotides)

TTTATAGGAC TICTAATCTA ATTINCCTAT AGIGIGACTA AAAGGGAGGC AAATTATIGG AACGGATTAT TCAAATGGNT CCTTAAATAT TGCTATGTAT AATAAGCCAG TTATTATATC AGGACCATGT TCTCTGTAGG CCACTGTTTT NCTCTCTCAT TCTCCAGTGG CGGCGGGG GAAGGCGGAG GCAGAGGCAG CAGCAGCCGC GCTGGCTGCA AATGAATGAN CCCCCAGCTT GGGGGGAGGA CTCCAGGTGA GCCTCTGCCC TCGGGAGGCCC C

SEO ID NO:1499: (Length of Sequence = 395 Nucleotides)

TITIATCACA CCCIGITITC CAAGGGICCI GITACGIACC ATTCACCATT CTGCTTAGCA ATGGCTTGIG AGATGGCATT
TATTCCTTCA GCATGIATIT INATGITCAC CITCCICTCA CCTAAAITCC TCCCCCACCC CAATAACAAT TAGITGITCT
ATTTGCATGI AGCCAGAGCA AAAAATGATT TCTTTCCCTT AAGITACTAT TATTATAAAA GGGACGATAA ACACATGAGT
CATTATACCA CAAGIATAGT GIGGAAAGGA CTCTAAACAT AGGCTCACTG AAGAAGGIGG CATTTGGGCC AGGGCTCAAA
ATAAGGCAGA TTCAGATTTG AACTGAATAG ATGGAGGAGT CATTTCAAAC AGAAGGAATG NCATAACATG TGGAG

SEO ID NO:1500: (Length of Sequence = 272 Nucleotides)

CTGAGTAAGN GITCCCAGIC GGICCCACTG GICACAAAIT TINIGGCACC GATCATIGAC ATTCACAGGG TOGTGATAGI CCAGITCATT GAGCICCIGC GCGATGGCIG CGATCIGCIC CACGCGGICC TGGIGCGCIG CCAGGTCGCI CTCGAACGNC TCGIGCTICC GCAGCAGAGC CCGNACCTCI NINAGCGACG CCGACTCGIA ATCCTINIGC AGCAAGATCI GCTCTITGCC ATAAGCCCAA GICTCGIGCG TIGAGGCCTI CT

SEO ID NO:1501: (Length of Sequence = 394 Nucleotides)

TITITITICC TEGACCIGIC ACAAGCITIA TIGICCOGAG CACAGACICG CCACACITCA ACAATICCAC TGIGGGGAGG
GGAGGGGIGA ATGAAGGACC TGGGGAGGG ACATGGCIGA GCCACANCCG GGCGGCCACA CGGGGGGGC TGAGAGGGCCC
ACGGAGGCAG AAGCICCCAA GGAAACCGCI TCTTGGACAC CCGTCACCAG GAGCCCACCI CCGGGGGCIC AGNICCICCC
GGCACCCICC TAGATGGACC TCTGGCIGIT AGIAGACTAA TCGGTGCCCC TACCGATGGG GCAGAGCTGC CTGATTITIG
CTAGAAAGAG CTGTATTTGA NCCINGGITA GGNCACTAAA GCATCGTTCT AGACGGCTGT TAATAGAACT NCAT

SEO ID NO:1502: (Length of Sequence = 373 Nucleotides)

GANACAAGGC ATAATGTTGT CACAGAATCA GAGATCCAGT CTCACTTTTC CACAAATCTC CAAATCTCCA GTCTTATCTT
GTGTGCTCTA ATGGTTTGGT TCAATCCCTT TCCAACTCTT GTTTTCAAAG CATGGGGCCT GAGTGTTCTC CACTCCTCCT
AAGAAAGGAG CTTGGGTGGA AGGGACCATG CTGACCTCCT CCATCAGAGG GCTCTTCCAG TAGTATCTC GGATGCAACC
TCCATTTCTC AGTTACCATT ATTTCCTGTA TCAGCTTTGT CCTTCCTGGN GGGATGCACA GTGATCCGGG CCACCACTGT
TGTTGTCTTG TGCTTCTGCT CTTTCCTATG GTTTCAGGNT ATTTTCTGGG GTT

SEC ID NO:1503: (Length of Sequence = 265 Nucleotides)

GNCAACAGGC CAGINITTAA AGAGGGTCAA GTGGAGGTGC ATATTCCAGA GAATGCTCCC GTAGGTACCT CTGTAATTCA GCTCCATGCC ACTGATGCAG ATATAGGCAG TAATGCTGAA ATCCGGTACA TTTTTGGTGC CCAGGTCGCC CCTGCAACCA AAAGACTCTT TGCTTTAAAT AATACTACTG GGCTGATTAC ANTTCAGAGG TCCTNNGATA GAGAGGAGAC AGCCATTCAC AAAGTGNCAG TGCTGGCTAG TGACGG

SEO ID NO:1504: (Length of Sequence = 311 Nucleotides)

SEQ ID NO:1505: (Length of Sequence = 363 Nucleotides)

CCACTCATGG CAGAAGGGAA GGGGAGCTAG TGTGTGCAGA AATTGTATGG TGAGAGAGAA GAAACAAGAG AGAGGGAAGG
GAGATAGCAG GCTCTTTTCA ACAACCAGCT CTCATGGGAA ATCATAGAGT GAGAACTCAT TCACTACCAT GAGAATGGCA
CTAGGCCATT AATGAGGGAT TCGCCCCTAT GACCCAAATA CCTCCCATTA AGCTCTACCT CCAACACTGG AGATCACACA
TCANCATAAA ATTTGGAGGG GTCGAATATC CAAACCNTAG CAACTTGGAA CCACCAGAAG CTGGAAGAGG CAAGGAAAGA
TTTTNTCCTA GAGGCTTCAG AATAAGGTAT TGCAATTCTG AAA

SEO ID NO:1506: (Length of Sequence = 177 Nucleotides)

CGGACAGAGC AGGGCAGAAA AATGAGGGAA GGATGACAGA AGCTCATCAG AAAGCCAGTA ATACATAAGA TTAGTTTINT
CAGCAAAACC TNGTAAACTT TGACGTTAAA AGACAAATAT TTTGATCTCT CATTCCCACT CTCAAAAAAGG TTTCTAGTTC
ATATTGTTTT GCTAAAA

SEO ID NO:1507: (Length of Sequence = 345 Nucleotides)

CTTGCTTGAT TTTCCCCTGT GTGTCAGAGA ATGTGCACAT TGAAAGAGAG GGAGCTCTCC ATCACCAAGA GAGCCCAAAA ATAGCCCAAC TGATCATAGC CGTTGTAAAA ATATTCATGG ATGTAAGGAA AGATCCTTTC CCAGTCTGAT GCTCCTTGAC TTGTGATTTG CTAAATTTGA GAAGCCATCA CTTACACAAC CTGTTTTATA GACAAATCCT TCCAGTTTCA GAAGAAAAAA TGTCATCTAT CTCLACCTCC ATCTCTTTTT CAAACTTCGA TAGATGAGAA GAAAATGGTG AAATAAATTT TTTAGAATCA GTTTTGCAAG ATTGGGTTTC AAGGA

SEO ID NO:1508: (Length of Sequence = 326 Nucleotides)

AGTTGGATTT CAGCIACTCA GAGTAATTGG AAAAGGCCAC AGCCTGGTGG GCTTCACAGC TTTCAGAGAC CTGGTAGGGG
ATGGCTAACA GGTTCINCTG CCAGGAGACA AGTGGCAGAC CCAGGTGTGA AACTTTTACA GGTCCCACCA AGCCTTTCTT
ATGGAGCACA GAGCATAAGG ACAACTTCTG CAGAAATGGA ATGGGGTACT TGGAACCAAA AATACATACA CCTCCTTTCC
CACCTGCCTC CAGCTTAGTA GCCCATAGTC CTCTTTGTCC CTCACACTGA GCCAGGGCCT GNCTTAGATG ATGAAATGCA
TGGCCT

SEO ID NO:1509: (Length of Sequence = 329 Nucleotides)

AGTATGGGTC CCTTGGTACT ACTCAAGGTT TACAATATTG CATTAAACAC ATTGAAAAAT ACACGAGAAC CTTGAGGGAT CACATTTTAC TGCAATATGT GATTTCCTGG TGAGGACTCCT TGTGCAGAGA TGATTAGCTC ACAGAGCGTT GTAAGCACGT ACTCGCAACA CCTGAGCATG CCGCAATGGC AACAGGAGGT ATCTTCACAA TTATGATGGT AGTACAGTAT GTACTGCAGT TGTTTACACA GTTATGATTT AGTACTACAT CTTTTACANIT GCNTATTTNC TINCTATTTT GAATGGTATG TACTGTCTGT GTGTACATA

351

SEO ID NO:1510: (Length of Sequence = 247 Nucleotides)

TAGGAAAGAG TAAGANCTTC TTTNCAGGCT GGAGGTGCTC GTATGGTGGG ACAGGAAAGG GGAAAAGAGA AAGGGGCAAC ATGGCAGACA TACCACGGTT CCTACAGAGA TTAGGGGCAG CCCTGGCCCG GGAAGTACAC AGGGCAGAGA GCTGACTCTC AGGCCAGGAA GGAGTTTAGC TCTNACCCAT CCTCANGGAC CACGGCTCTC CCCCAGCCTC AGCTGACACA CACACAAAGG AGGGTTTT

SEO ID NO:1511: (Length of Sequence = 369 Nucleotides)

CCACTTGCTC CTITATTAAC TGINCITCCT GIAGIGIGIA TITGGGATCC ACIGGGAATC ATAGAAAGGA ATCAGIGCTA
GENICIGITG GGATTGCACC CTGAGGGATG TGGCTITGGC TICTCTATCA ACCITTCTGT TCCCITGTGC TATAGGAGIT
AAGTCCCTTT NATGCCCCCT ACAGIGGATT ATAGCTATGG CCTGTGGCAG GIGTATTGTT TACAATAGCT GAAGAATTTC
AGGCCCATGC TITATGGGGG AGGGTTTINC TAGCTAGTAG TCCCCTTTCT TTCTAGATTG CAGCATAAGC GTGAACCNCC
AAGGAATGCC ATATTTTAGA ATCCTGTNAT AGGATGGTTA AGGCTTTTT

SEO ID NO:1512: (Length of Sequence = 236 Nucleotides)

ATGCATTAAG AAAAGACAGC CAAATGACAG ACTGATAAAA TATTTTCATT ACAAAATTGG TTGAGAACTA CCGTGTGACG
TAAATGAAGT TICTATTACA CATGTACTAA CAGAGACTIT TCATTACATA TTCTAGGATA TATTTAAAAT ATATGTATAT
TTTGATATTA AGGGAATATA TTTTGTTGTC ATTTTACAAT GTGTAACTAC ATATATATTA NGGCCTTTCC AATAAA

SEQ ID NO:1513: (Length of Sequence = 408 Nucleotides)

CATTAATATT CTCAGTGTTG GAAATATTTT NATATTGCCA AGACCATAAT GTGAGGRGTG CAGCTGCATA ANTOCCTGAG
AGAAGATTAG TGGGGCTAGC ACCITACAAG GAAAGACAAG CTTGTTGGCT GGGCCCAAGG ACAGTCAAAT GTCTGCCTGA
CAATCTCCAC ACAGAAGGGT TGCTCAGATC ACTTAGGACA CCCAGAAAGA GCTCACAAAG GGCAAACAAC CTAAGGCTGN
TATTCTCCAT CTAGCGGTAC TTACCTGGGA ACTGAGTGGC AGTGGACAGG AAGCAGGGCC TGGGCTAGGG AGACCCTCAG
GAGGAANGGG GACCCAAGAA GTTAGAAGTC CATTCATTCA TATACTCATT CATTCAGCAA ACATGCGCTT GACACCTTCT
GTTATGCT

SEQ ID NO:1514: (Length of Sequence = 359 Nucleotides)

TINNOCAGGO TEGTOTOAAA CTOCTEGECT CAAGINATOO GIOCACCTTE GOTTOCOAAA GINCTAGGAT TACAGGOATE
AGCCACTGIN COTEGOTAGA AAAININITT TIAAAAGINA GGATGIAGAA TINOCTAGCT ATGIAGGCAA GGCAGGAGGA
GAGGGGCCCA GITGGGAAGC ATAGCCCACA AGAGTATGAG GGCCTGANOO AGGATGGTGG CAACAGGGAT GGAGAGGAAG
GCGTGCCAGG GCATGGTGGC TCACACCTTA TAATCCTAGC ACTTTGAGAG GCTGAGGGAG GAGGATCATT TINAGCCCAA
AAGTTAGAGA CCAGCCTGGG GNAACATAGT TAAGGACAC

SEO ID NO:1515: (Length of Sequence = 343 Nucleotides)

GAGCCCCTTG ATGGCAAGAN CTGACCCTTC CATCCTGGAG AAGAGGAGAC CAATTINATA TIATGGAGGC AGAATATACA
GGACTGTGTG ACTAATTCGA CATGTGTGTC CATGGAGCTT GAAGGGGACA GAACCACAGG TGCAAAACTG GTGTAGGTAG
TGCTGGCCCAT TGCTCAGAAC TTTGTGTGAG TTGAGCCCCAG GCCTCTGGTT GCAGGACTCG TGAATGGAGC AGTTCTGAGA
ACCACCCTTT TGCTAAGGGA GCTTNGGAGC CACATGGCTG CTCCCTTCAC ACTGGGTAAC AGTGTAGTAT CCTGTGAGAG
AATAAACGTA TTCATTTAAA AAG

SEO ID NO:1516: (Length of Sequence = 380 Nucleotides)

TTTTGCCTTA TTCTATCCGA TTTTTTCCCT AAGCTTCTAC CTGGNATTIN CCTTTGGAAA AGTCTCTGAG GTTCCACCAA
AATATGGAAC TINATTTTGG ACACTTTGAC GAAAGAGATA AGACATCCAG GAACATGCGA GGCTCCCGGA TGAATGGNIT
GCCTAGCCCC ACTCACAGGG CCCCACTGTAG CTTCTACCGA ACCAGAACCT TGCAGGCACT GAGGTAATGA GAAGAAAGCC
AAGAAGGTAC GTTTCTACCG CAATGGGGAC CGCTACTTCA AGGGGATTGT GTACGCTGTG TCCTCTGACC GTTTTCGCAG
CTTTNACGCC TTGCTGGCTG ANCTGACGNG ATCTCTNTCT GACAACATCA ACCTGCCTCA

SEO ID NO:1517: (Length of Sequence = 411 Nucleotides)

TGAGCAAAAC ACAGAGGACT GCACCTCTAG TGGCTCGTAA TGAGAAAGAA GATGGTCTCA AACCTGAGAA AGATAATGTG
GAGTGGACCT CTGTTGTCTC AGTATTAACA GTCCCTTCTA GGAAGTAGGT AGCATTTCTG AAAATAGAGT GAAGCAATTG
ACTGATGGAT TTAATCTTTA AACTGCTTAG GTAACCATCA ATCTGTAATG AGCTTAATAC TCTTAACTAG GTGCTATTTT
NCATGTGGC TACTTTGCCA GTGATAAAGG ATTACGAAAA ATTCTTTACC AGAGGAAAAA AAAAAATTGA ATGACCTTTC
TTGGGAAGGT GGTCCCTTGT TTGTGATCAA ACTTTGACAA GAACTGGTAA TTAATTTCCT CTAAGGAATT NACCGTTCTC
ATAGTGTGTT T

SEO ID NO:1518: (Length of Sequence = 388 Nucleotides)

GGIGGGCAGC TICTCTCTGC AGCTGCTCTC CCATCATCTG GCTGAATATG GGGCTTTNAT GGGCCTCAGG GGAGGAAGTG
TGTGCNAAAT GGTCCGTGGG CAAACATGGG CGGGCCTGGA AAAGGCACCA CAAGTTCCCA CCCCAGTCAG TAGGATCAGC
AGTCTGACAC CCAGGCTTCA GGCCCTCCCC GACTTGAAGG TGGTGCTTCA CCAAGGACTC ACCCACTCCT GCCCAGGAGC
TTGINTGCCT CCTGCTGCCA TTTATGGTGC CCAGGCTGTT TGINCCAAGG AGTGTCTGTG GGCCAGGNCT GAGCTGCCCT
CAGCACCCCC TTGGCCTCTT TTCTGTNCTC ATTGGTGCCC AAAGTCCGCA GCAGGCTGAA GTGGCAGG

SEO ID NO:1519: (Length of Sequence = 358 Nucleotides)

TTGGITAAGA CCAAAGICAG ATCACICCCI CCTAGCICCA AACCTGCAGI GGCICCCAAT TCINTCAGCA TACAAACCCA
GATCCTCAGG CTGCCATTIN TGGGCTGAAT CCTGICCCTG CTGTCTGATC CCACCAGACA TAATGGAGGC CTGAGGTTCC
CTGAACACTC CTAGTTTAGC CTTAAGTTAA GTATTTGCAC ATGCTGGTTC CTATGCCTGA GATAATGTTC CACATTINAT
CCCATTGCTT GCCAGAAATA GAAACCCTTC CACATAATTN CAAAACAGAG TTTACANCAC AGAGCTTTGG GTGACTGCAG
GCCTCCAAGA ANGGNAGGCA GAAGGGGCAC TGAAGAGT

SEO ID NO:1520: (Length of Sequence = 379 Nucleotides)

CCAGAGITAA ATATGCCCAG GCTGAAAGAA GGTGTATAAT GTATGGNCGT NCTTATACCA AATGATTTCT TTGGAATTTA
AACAAATATG TTTAGTATTT TATTCCTAAT TTAGGAAGAA AAAGCAACTA AAGTTGTNCT GACATTGTAC ACAGATGAGT
AGCACGTAAC TTTTATTTAG TAAGCCCCAT AGGATAGTAN GGNATAAAAG TTGTTAGTGA GCAAAACAGG AGTATCCTGC
CATTTGCTTT AATTCTNCTT GTGATAGTTT TGAGGGTACA ATAATTCCTG TGTGCGTGTC ACTCAAGCAA ACCAGAAAGT
GTCTTTTGTA AATACGCATT TTGGGCCTCA TCCTCATGGA GGTTCCCGTT GTTTGTTAGG

SEO ID NO:1521: (Length of Sequence = 339 Nucleotides)

GGGACAGGAA GCCTCTTGGG TTGGACTCAG ACTCAGGAGG TGACTCAAGC CTCAAGCTCA GAAGCCCTCT GINACCATCT
GTTGACTCAG AAGCATGCCC ACCATCCCAT GCAGTGCCCT TCCAGGCACT GTCCTGTAGC AGACGGAGTT CAGGCTTTGG
AAGTAGACAG ACCTGGGTTC AAATCACAGC TCCGCTTCTT CCGCCTGAAG CTCCATAACC TAGGATAAAG TCGCTAAGCC
TNCCCAAGTC TCAGATTTCT TACCTCTAAG GTGAANGGAT TGGATTCCAC TTTACTTCCC CCCTTTTCCC TTTANAGGACT
CTGCATCCTC NTTTGCTTG

SEO ID NO:1522: (Length of Sequence = 405 Nucleotides)

GIGAATITCA AGCAATIGIT AATGGGGACC AACAGGGCTG CATTAAGAAA ACCACTIUN ACTGATCTCT CCCCCACATA
TITTIAATIT GICTIGCITT GITTATITTG GITATGCAAG TCCTTTCTCT TCATGAAACA AGIGTAAGGC TCTAAGGCTA
AAATAATAGT TATTITTGIG GGCCCCAAAT AGCTACTTTT GAATTTCTTT CTTTAGTATA TCTCAAATCT GGGGAACATG
GAACTIGAAG ACTCCTAACC ATGAAGCATT TGGAAAAATA CATATCATTC ACTTTTCACA GAACCATTTT CTTAAAAATA
AGGGGGCCAAT ATCCAGATTC ACATGCATGT TCATAAATAA AGCTTTGGTT TTAAAACAAA TCCACACCAG CAATTATTTT
CAGCT

SEO ID NO:1523: (Length of Sequence = 284 Nucleotides)

AGNICACAGA ACTOCAATIC TITATTAATO ACAGCTIGOT CACAATGACA TACAGGAAAA TAGCACTAAT GAAGNGTAAA TATGCAGGCA GCAACCTICA GGAGTIGGGA GITGGGGAGA AACENCITCA AAACTGCGAT AGGTACTTAT GGTGGGTATO TGGTGATTOT NAGITGGCAC AAATGCCCTG CCTAGCCCCC TTAACTGCGT CACTITCACA GATGCNGTGT TTTGTTGTTG GTGTTGTTAG TAGGCAGGAT TGCCTTACAC TGGGGAAGAA AGAC

SEO ID NO:1524: (Length of Sequence = 299 Nucleotides)

GIGCTIGIAC GIGACAGITT IGICIGATCA CATITIAGGA AGAIGAIGCI GITCIINCIT CITAAGIATI TATITINATC
AGICAAGIGA TAGGAAGITC AATTICAAGI ACAAGACATI IGGAICAAGA AGIGACIATI ATTIATITAT TINAGAIGGA
GICTIGCICI GITGCCCCAAG CIGGAGIGCA GIGGIGIGAT CICAGCICAC IGCAACITCC ICCICCIGGG TICAAGCAAT
TCCNCIGCCI CAGACICCCG AGIAGCIGGA ATTIACAGGC ACCCACCGGG ACCAGIGAA

SEQ ID NO:1525: (Length of Sequence = 398 Nucleotides)

GCCCATGAAG CAGCTCTCGT GGATTGGAGT CTCATGCCTG CAGCTCTCCC ATACTGGAGT TGCATGCTGG TGGTCCTACCA
GTGCTGGTGT CTGGGCAGTG GCCTCACTCC CATGGCTCCA GGAGGCATTG CCCTGGTGAG GGATCTCTGT GGTGGCTCTG
TCCCTGTNAC AAGTTTCTGC CTGGGCTTCC AGGCTGTCCA TGATATCCTT TGAAATCTAA TTGGAGGCTG GCATGACCCC
ATGGCTTCCA CACTCTGTGC ACCTGCAGAA TCAGCACCAT GTGGACACTG CCAAGACCTA CCTACCACTT GTGCTCTCTG
GAGCAGCAGC ACAAGCTACA TCTGGGGCTG CTTGAGCCCAT GGCTGGGGCT NCCAAGGAGC AGAGTCCTGA GGGTGGCC

SEO ID NO:1526: (Length of Sequence = 318 Nucleotides)

GICTCTCTCT ACTGCACCAT GATGCCTTTA AAAAGAATCT AGGGGCTGGG CACAGTGGCT CACGCCTNTA ACCCAGCACT TTGGGAGGAG TTCACTTGAG CTCAGGAGCT CGAGACCAGC CTAGGCAACA TAGTGAGGAC CCCGTNTCCA CTAAAAATGA AAGCAAAATTA GCTGGGTATG GTGGTCCATG TCTGTACTGT GGTCTAAGCT ACTCGGGAAG TTGAAGCAGG AGGNTCACTT GAGCCCAGAA GGTCAAGGCT GTAGTGAGCC ATGATTNTGC CACTGCATTC CAGCCTGGGC AACACAGTNA GACCCTGT

SEO ID NO:1527: (Length of Sequence = 313 Nucleotides)

TTGGCTAGAA GGGAGGCTGG AGCCTTTCAT GGTGGCTTTT GAATGCCATG GTGAATAGTT TGTCCTTTAT TTGTNATTGA
ATAGCAATTT GTACACTTCT GAGCTATTAG AGTGAAATGA TTAAGCCTGT GGTTTAGGAA GAAAGAGCCT ATTAGGGAGA
TAAATCTTTC CCTAGTTGTA GGAAGGGTTG GAACAGTATG ATATGGAGAG GGTAGTAATG AATGANGGAA TNGAAAAACGA
GAATAATTTC AATGATACTG GAGGTGCAGT ATACAAGTTG NGCAGTAGGT TTATGTCTAG GAAGATAAGA AGT

SEO ID NO:1528: (Length of Sequence = 405 Mucleotides)

GCCGTCGCTA CCGCCACCGC CACCGCCACC GCCGCCGAGT GCTGTCTCTA TGGCGAGGAG GAGGAGGAGG AGCGCGAGTC AGCGACACAA GTACATAAAT AAAGGATAAA ATATTTTATG AAACAAATCT TCAATCAAGT ATAACATTTT GATGCTTGGC

ATCTAGACTO COTTGTGCCO TOACTATGCO AGOGGAACTG TAGATCATAG COAAAGAATT TGTGAAGTTT GGGCTTGCAA CTTGGATGAA GAGATGAAGA AAATTCGTOA AGTTATCOGA AAATATAATT ACGTTGCTAT GGACACOGAG TTTCCAGGTG TGGTTGCAAG ACCCATTGGA GAATTCAGGA GCAATNCTGA CTATCAATAC CAACTATTTC GGTGTAATGT AGACTTGTTA AAGAT

SEO ID NO:1529: (Length of Sequence = 241 Mucleotides)

GAAGGAGAAA CACTICITGC CICCATAAIT CAGACAGIAA ACIGATCGCI GAGATIGAAG TITGCITGIT TCCIGGGGAA
GCTINAAGAT CCICGIGGGA CCACCATCCC CIGCICAGIC CICCCIGGAA GGGGGCACIG GCIGGGTATG AGCCGCGTCA
CCGITGGGIT TGIAACITIN TGGATGGIGC CIGGNITICA CCIGGGGCIG GCIGAGGAAA GGGGAGGCGG TAGGNGICIG
C

SEQ ID NO:1530: (Length of Sequence = 356 Nucleotides)

GGCCCCTCCA CACCTAGAGC GGTGCAGGAG AGGACT

SEQ ID NO:1531: (Length of Sequence = 379 Nucleotides)

CCAACAGATG CIGCTACGIT TCCTTCAAAA TIGITAAACA TCTCTIGCGG AAGAAGCTGC TTAGITATAT CCAGCGATTG
GTTCAAATCC ACGITGATAC AATGAAGGGT GGGGTATCTA GCAGGATGIC TAGITCACGC ACTGGGTGAA AAACAACCAG
AGCTGCAGAT AAGTGAACGA GATGTTCTCT GTGTTCAGAT TGCTGGACTT TGTCATGATC TCGGTCATGG GCCATTTTCT
CACATGTTTG ATGGACGATT TTATTCCACT TGCTCGCCCG GAGGTGAAAT GGACGCATGA ACAAGGCTCA GTTATGATGT
TTGAGCACCT TATTTAATTC TAATGGGATT AAGCCTGTCA TGGAACAATA TGGGTCTCA

SEO ID NO:1532: (Length of Sequence = 307 Nucleotides)

GATAAACTIG AGCCACCAAG AAGIGGACIC IGCCIAGGAA GACAGITIGC IGAAGITAGA AAGIACIGGI CIAGGAACCA GAAAACCIGA ITCINCCCAA GAGITAGAAT IGINAGINAG ITCIINCIGG ITTINAGIIT CCIITATCIGI AAAATAATTA CCCAGITCAA ITGGATAATC ICTATGATCC CIITCCACATI CIGCATACTI GGATATCIAC IGITICIAAA TATIITIGGCA ITTCIITATAA AGCCCIITCA CATTINCIIT ATTATIITIC CCICACAAGA ATTCCIGAAA TAGGATA

SEO ID NO:1533: (Length of Sequence = 337 Nucleotides)

ATGCCTTTAT TIGCIGATIG AGAAGIGGIC CAGCCGIGGG CTAGCAGICA TITACATATC AGIGACCAAA TGCAAACATA CCCGIACIAA CAGIGCITIG GICCATGACA TACCCTTITIG ACAGCCCAAA GCTGAAACGI CAACTCIATC TGGGGITACT TGCTTATACA AAGATGITAC TCTAGCAATT GITGCTTGAG GGCAAGACCN GATGATTGIC ACTAGTAGGA AGAAAGCAGA AGIGATGCAG CTTACACTGC ATAGTCCCTA CCCTINTGGA TTAAATGGAA AAGITGCTCA AACATAAACT TGTTCTTAAC AAAGGTGGGT AAGANTC

SEO ID NO:1534: (Length of Sequence = 317 Nucleotides)

ATGGGCATGT GGGTACTACG TTTAAATATT TAATTATTT AAAAATAAAA TAGGAAAGAT AAAATAGCTT AAAGTGTATT GATGCTCTGA ATAACTTTAT GAGTGAATAG ATACTGAAAT TTGAAGTCAG TGTTTTGCAC AACAAATCAA GATTTGGGAC TGGACTTACT GGGTTGGGGA CTTCTTAGGG ATAACGGTGG TGCTATGAGC ATGCTGGAAA GATGAGAAGC AAAAGCCTGG AATTGGGAGT CCTGTACTGT CTTTAGGGTA TGCAAAGAGG CTCCTTCTTT TCTAGGTGTT CATCAGTACA ATATGAC

SEQ ID NO:1535: (Length of Sequence = 323 Nucleotides)

ATATTACATT GATGICAGIC TITAAAGATG GAGTAGGACT TINCAGGCAG CAACGAAAGG GAAGGACATT TCAGAAGCAG
AAATACCATT TGITAAGGGA TGACAGCCAA GAAATATTAA AGCATATTIG GAAAGTATTG AAAATCICIG TGIGGCTAGA
ACITTAGATG AAGAATCAGA TACATCIGGA GAAGGAGATT NAACCNGATG ATCATAAAGA ACATTTIATT TAGGCCATGG
TAAGGCTIGG GCACINIGGA GCCCATGAAG GITTITGGAC AAGGGAGTIT CCTTAGGGAG GAGTATNAAG CCATAAACCA
AAT

SEQ ID NO:1536: (Length of Sequence = 305 Nucleotides)

AACCACATIT TIACIGCATC INCICCACGC TGGATTCCAA CATGCTGGCC CGGAGCGTGG CTGGCTGGAA GCAACTCCAA CAGGITITIC CCITCCCCGT CATGIACATT ATTIATITIT GATCCTACTC ACTGTCCCAA GTCCAGAGGC AGITACAAAA AACACTCTTG ATGCAAACCG TGAGIGGCTA CAACACACGG ATGGGGGTGG GCGCGATTCC CACAACAGGG AGTGGAATCC GGGGAAGATG ATATATAGGG GCAAGACGGC CCCTTACTTT GCTAAGAGTA TATGGGAGCT CAAAA

SEO ID NO:1537: (Length of Sequence = 279 Nucleotides)

GGTGGCAGCG GCGGCGCGC GACTGAAGCG CGCGAAAAGC TGAGGCGGCA ACGTCGGGGA CGCCTGCNCG GGACGGCTCT GTAGGAAGGA ACTTGGTTCC CCCTCCCTCA GCTTCCGCCC CAAAAGATTC AGAATGGACA GTTTAGAAGA ACCTCAGAAA AAAGTCTTTA AGGCTCGAAA AACGATGAGA GTNAGTNATC GTCAGCAACT TGAAGCAGTG TACAAGGTCA AAGAAGAACT NTTGAAAACT TGATGTCAAG CTGTTAAATN GCAACCATG

SEQ ID NO:1538: (Length of Sequence = 310 Nucleotides)

ATATTICCIT CIGCICIGAC TCCGGAAGAA CITGCACIGI TGCCTAGGCI GATAATCCCC GAAAAAAAGI AACAAATGCA
ATINI'ACCCC CCACCCCCAT ATACAGCCCI CATATATATA TATGAGAGAG AGAGAGGAAA AGATCATGAG ACATGICITC
TAGGGAAAAA AAATTCTAAC TTCCCTAGCC ACTGTAGTCA TTTGAAACCI GAGTTAGACT ATGAGTTAGG AAGTATTTTC
ATAGAGTTCA ATTAATATAT TTCTGCTCTA TGCATGGATG CTAACAGGTT TAAGGAAACA CAAAAGCCAA

SEQ ID NO:1539: (Length of Sequence = 267 Nucleotides)

GAGATITIAC TITGIAATOG AGIAATITIAG CCACACTCIT GIGAGGGAAC AAGCCAGAGC CAGGACCGCA TATIACCOGG TAAAGCTGCA GAGAAGACTT GAGACTIGIA AGATTGCNCC NGGCTGCAGT CCCGTGGTCA GIAACATCTG CAACATTATA CAGCCAGCAG ATCAGCTCIT CCAGCTGACA GCAAAATGTC TTCACACATT GCACCAGTGA TTCTTTTCCC TGINCTCCTC CTTTCCTGGG GAAGCTGCCC TTNAACA

SEQ ID NO:1540: (Length of Sequence = 354 Nucleotides)

ATTTATTCAG ATGAAAAAA ATCAAGGCTT AATTTAAGTA ACTTGTCCAA GGTCAAGGAG TTGACAAGTG GCTGAGCTGG
AGTTCAGCAT CTCAGACATC TTCCTTTGAA TCCTTGCCTT CCTTGTGAAT TTCAGATGAC GGAGCATGAC GGCTGCATGA
TTATGGGGTC ACCGGGCCTG TCCTGGGCCT GAGGGACCAA GGATCAGAAA GGGCAAGAAC CAACTCGNTC AGCTAGTGAA
AGTGCAATTG GACANTGATC CTGTTTCCGG GNTTAACCTT CCGCTTGGCC TTTAAGAGGG NITCTTGAAA TGCACCAAGG
GGGCCTAGAG GAAGCAAGCA AACTNCTTGG ACCT

SEO ID NO:1541: (Length of Sequence = 403 Nucleotides)

GIGATGITAT ATCAGGIAAA ACCTGTCTAA GGAGAATAGA CAGTAGTTAG TICAACTTAC TCATTACGTA TTAGGAAGAT
TAACCIGGIT ATCATTGITT TATACATATA TATATGNAAT ATATATGAGT ATTCGTATAA ATATAATACT TTTACCITGI
TTATGIATIT ACTCAATATT CICCITTICC TCTAAAATAA TCTGAAGIGA CTATTATCAA TAAGITTACT ATGCCAAAAT
TCATTAATTG CCTTTCACTT AACTITIGGG GCCATAATAA ATAATAAAAT GTATTGCCAT AACATTAATA AACTACCITA
CAAAACCACC AATTAAAATC AAACAACCAA AAAGGIGITA TTTACATCIG NNCACATAAA TCTACTAAAA ATACAGGGIT
CAT

SEO ID NO:1542: (Length of Sequence = 333 Nucleotides)

CTGGTACATG ANTITATAAA AACATGTCAC GCCCGGCTCT GTGGCTCATG CCTGTAATCC CAGCACTTTG GAAGGCCGAG
GCGGCGGNT CACAAGGTCA GGAGATCGAG ACCATCCTGG CTAAAACGGT GAAACCCGTC TCTACTAAAA ATACAAAAAA
TTAGCCGGGC GTGGTGGTGG GCGCCTGTAG TCCCAGCTAC TCTGGAAGCT GAGGCAGGAG AATGGCATGA ACCCNGAAGG
CGGAGTTTTC AGTGAGCAGA GATCATGCCA CTGCACTNCA GCCTGGGTGA CAGAGCAGAG CGGGGACTCC GGAGCAATGG
GNAGTACAAT CCT

SEO ID NO:1543: (Length of Sequence = 329 Nucleotides)

CCCCIGATAA ACCIATCAGA TICIGIGAGA CITATICATI GICATTAAGA ATAGCAGGG AAAGACIGGC CCCCATGATT
CAATTACCIC CCCCIGCATC CITCCCACAA CAIGIGGGAA TIGIGGGAGA TACAATICAA GIIGAAATIT GGGAGGCGGC
ACAGCIGAAC CATATCAGIC TGIATIATCI CICCNITIIT CIGCIITAAG NGACIATACG NAGGIGTIGI TITCAGGGNI
TATACATAGG TATICIGAAA GAIGGGGITA TITICIGIIT CANACITIGA CIAAGIGGCI TCITITGICC CCIATGIGCC
AGAATAGCC

SEO ID NO:1544: (Length of Sequence = 313 Nucleotides)

CGGAGATCCG TGATGTAACA AGGATTGANC GAATCGGTGC CCACTCCCAC ATCCGGGGAC TGGGGCTGGA CGATGCCTTG
GAGCCTCGGC AGGCTTCGCA AGGCATGGTG GGTCANCTGG CGGCACGGCG GGCGGCTGGC GTGGTGCTGG AGATGATCCG
GGAAGGGAAG ATTGCCGGTC GGGCAGTCCT TATTGCTGGC CAGCCGGGCA CGGGGAAGAC GGCCATCGCC ATGGGCATGG
CGCAGGCCCT NGGCCCTGAC ACGCCATTCA CAGCCATCGC CGGCAGTNAA ATCTTCTCCC TGGAGATGAG CAA

SEO ID NO:1545: (Length of Sequence = 384 Nucleotides)

CCCAAAACCT GGAGETAAGA ACTICATCIC ACTITIGACA CCCCAGCCCC CAAAATATGG AAGCCCAGGA GAGCCAGGAG
AATTIATAGC AGAGGCITAA AGAGAAAGIT ATGATTIGIT TAAAGIAGAG AATAAGGIGA AAAATAAAAC CIGGTACTCT
GTCIGGAAGI CCTGGAAGIC TCCTIGCCCA ACCTCAACIG GCCTGIGGGC TCCTGINICC TIGCTCTGGG ATGCCATGGT
GAATGIGAAA ACAGGGGAGG TIGIGIGIGG GGGTGGGAAT GGCCINICGG TIGCAAGGCG AGICCTITGC TGAGCCCAGC
CTGAGACCCA GCTTATGGGC TITATCCAGG TGAGAAAATN CTGGGGACAT GTGITCGAGG TTTA

SEO ID NO:1546: (Length of Sequence = 345 Nucleotides)

TTTAAAGAAC AATGATTAAG TGAAAATNCT CTCAGTTTT TTTAATTGGT TCAGCAATTG ATTAATTACT GAATCITGAC CCTAAACTTT TTAGTCTAGA AATGIGCTTG AGGAATACAG GCTGGAGATC AGCTTTTTGA CATTGCATTC CCCTCCTGGN TCACATCCAT GTTGGAATCA ATTTATAAAC TGCCTTCCTA AGGCTTAAAA TGATGGTGAT CTACAGACAA GTGCCTTCCT AGGCACAGGG TTGCTGGAGA CTGATGCCAG GCCCATGGCT CTTAAACGGA ACACTGAACT CATGGCAGAA ATGGTGGAAA GTAGAGAAAT GAATAGAGGG GGGAA

SEO ID NO:1547: (Length of Sequence = 342 Nucleotides)

GGAGGCTGAG GTGGGAGGNT CACTTGAGCC TGGGAGGTTG AGGCTGCAGT GAGCTGTGAC TGCACCACTA TACTACAGCC TGGGAGACAG AGTGAGACCC TGTCTCATAT ATATATATAT ATATGTATGT ATATATATAT ATGTTATATA ATGTTATATA ATGTTATATA TATCTAATAA ATGTTATATATA TATCTAATAA ATGTTATATATATA ATATATATATA TATCTAATAA ATGTTATATATATA TATCTAATAA ATGTTATATATATA TTATGTTATA TTATGCTATA TTTGGCTATT TCCTAAGAGC TCTATCGTAT TATTTCCATT TATTTGTGAG GA

SEO ID NO:1548: (Length of Sequence = 334 Nucleotides)

GGAAATAAAG GTGACATGAA CTAACTATTC AATCATGAAT GGTAGAAAAA AATGAAAATG TAACGAGATG GGATCCGGGT
CAAAGTCAGG GGAGGTATAG TTGAAGATAT TGAAGGAGTC ATTATGATAC CAAAGAAAAT GGAAAGAAGT GGTATCCAGA
TAGGTTATCC TTGGAGAGTA TCCNGGGATG TCTCTTTTCC TAAGACCTTA GAGAAGGAAA GGATGGCTGA TAATATAGGG
AAAAGTTGAC ATGGAAGGAT TAAATAATTT TTTTGAGGAA TTCACGTAAG GNATGATAAT CTGAATTTTC AGGGCTAGGC
TCAGAAGCAG GAAT

SEO ID NO:1549: (Length of Sequence = 362 Nucleotides)

SEO ID NO:1550: (Length of Sequence = 328 Nucleotides)

GGACTAATTA ACTAAAGAGG TITTGTACAG CAAAGAAAC TGTCAACAGA GTAAACAGAC CTACAGAATG GGAGAAAATA
TTCACAAACT ATGCACCCAA CAAAGCTCTA ATATCCAGAA TCTATAAGAA ACTTAAACCA TTGAACAACC AAAAAACAAA
CAACCCCATT AAAAGTGGAC AAAAGTCATG AACTGACACT TCTCAAAAAA AAGACATACA AGCAGCCAAC AAGCATATAA
AAAATGCTTG ATATCATTAA TTATCAGATG AATGCAAATC AAAACCACCC AAGTCTTTTT CTTCTGTCTA GGNTAATTTA
TTTTAGGG

SEO ID NO:1551: (Length of Sequence = 365 Nucleotides)

CAGGAATTIA CATGGGGAGA CCTACCTATG GCAGCTCTCG CCGTCGGGAT TACTATGACA GAGGATATGA TCGGGGCTAT
GATGATCGGG ACTACTATAG CAGATCATAC AGAGGAGGAG GTGGAGGAGG AGGAGGATGG AGACCTGCCC AAGACAGGGA
TCAGATTTAT AGAAGGCGGT CACCTTCTCC TTACTATAGT CGTGGAGGAT ACAGATCACG TTCCAGATCT CGATCATACT
CACCTCGTCG CTATTAAAAGC ATGAAGACTT TCTGAAACCT GCCCTAGAGC TGGGATATTG TTTGTGGGGC AATATTTTTN
ATTGTCTCTT GTTTAAAAAG TGAACAGTGC CTAGTGAAGT TACGT

SEO ID NO:1552: (Length of Sequence = 330 Nucleotides)

GATCCAAAAA AATTTACTGA AATAGCAAAA ACGTGGACTT TGGGATTTCC TCTAACTGCT GCAAATTATA ACACAGAATT
GCTCAGTGTT AATACTTGAN TTGTGGGGCC AAGTCTTCTG GCTGCCCTAG TTCTCTTTTC TGGCATTTGA AAGCCCTTGA
GCTAGCTATG GAGCTAATCT TTGGACAGGC TTTTTGGTTC CCAGGAATGT CATGCCTTTG AATTTCCAAT CTATATATAT
ACAGTGTGTG TGTAYGTATA NCTGTCTTTT CACTGTAAGG CACCTNCACC CATCCCTTAT AGAAGGNGGC CACAAACAAT
CAAGCAAATC

SEQ ID NO:1553: (Length of Sequence = 304 Nucleotides)

CCCTTGTCCC ACAGCCATT AAAAATCITC TGAAGGGCCT CAGGGCACAA AGTGATCATT TGGGATCCTA AGTTAAAAAG
GAAATGCAAG AGTAGGNTAC TCCAATTCCA GAGTCTTTGC AGGAGGCTAA TCCCACAAGA AGGGTAGCAT CAGAGAAGTG
GGCATTGGTC TTAGTGGTGG ATCATCAGGT AGACAAGTGA TAGTGTGTGT AACCCATCTG AAATTCATTT TACCGTCACC
ACTCTTACAA AGGACAGTTT ATTCCCAAGG ACAGTGCTGA CGGGGAGGGG GACAGGCAGG GAGT

<u>SEQ ID NO:1554:</u> (Length of Sequence = 309 Nucleotides)

TGTGTTACTG ACCATGTTT TGAGAGTAGT GCCCCTAACC ACTITGTCTC CACTTGCATA GTGTAGTGAT TTTNAGGNCT CTGTATGTCA TATTATAACA GAACTGACTG TATATGGCTA TTTTATCCCA TAATCAAGCC AATTCTTCCA GAATATTACC ATCAGTATTA CCACATACAT CCTCCCCAAAT CTTATTTCAA AGAATAAATA TATAGTCACT CATGGTTTTT AAGNAAACCC AAAACTACTC AACCAAAACC TTGAGGAAGG TTTTTCCAGG GNTTTCTACC TTAATTATTC ATAATGATT

SEO ID NO:1555: (Length of Sequence = 326 Nucleotides)

GTTTAAAAAC TGTCCAAATG TCATTTAAT TTATGAAGGC ACCCAGAATA AGINCTAATC TCATACTGCC CCAATATATT
TMCTGAAGCC AATTCTCTCT TTTATTAATT TTTACTGAAA ATAGCACTTT TTTCCTCCCC CTGATAGTAC TGGGTAATGT
TAGAAATGTCC TCTAAAAATTC TTTGGACCTT ATTTACATTC TCAAGAGNTT TTTTTAAATT TACCAATAAG ATGTGCTATT
TGAGGAATTA GACTTTAGTT CAGTTGTACA TGGNTTATGT CTGCTCATAT CATTCATGTC TGAGNCTTTC ATTTTATTAA
TATGGG

<u>SEO ID NO:1556:</u> (Length of Sequence = 375 Nucleotides)

CCCATCCCTG TITAGGIGCT TIGICCTCCT TGAGGAGCCT CCAATGCTGC TGCTCCTATA CATGTCACAA TITCAGACCC
AGCATGCTAG GAACTGCTGC CAGCGCCTGG TTAAGCCCAAT ACTAAATGGG GCCAAACAGG TGAACAGACA TTCTGTCTTT
CTCCAAACCT CTGAAAAAAGA TTCTGCAACT CATCTCACAG TAATTTGTTC CCTAATTTAC TCTTAGGAAA TTGTCGTTAA
AGTCTGATTA GGTTAAGTCC AATTCCCTGT AATTAGGATC CTCAGTGAAG AAAAATCTAC CCATCACCAC AATTTATTTT
CTTTTCTATA GCTCCAGCAT CAGTAATTGT ACCATTATTT TTGGCAGCTC TGGGG

SEO ID NO:1557: (Length of Sequence = 306 Nucleotides)

AATTCCGAAG ACTATTCCTA TACATTAGAG TGAATTINAG ACTATCTCCA TCATTCTCCA GCCATTCTTC AGTGGGAAAA AAACGGTGGA ATTAAACTAG TGGAAACAAG GCTTTCTCAT CTAGTCCCAA TCCAGTCGAT AAGCTGTGTT TNCCAATCAC TGCTCCAGCA CAATGGCCCT CAGTTTATTT TTAAGTCTAT GGCATGCCTG AAGGACCATG TTCCCATGAG TGACACCCTT CTGTAAATGT GGTGGCACAT TATGGGCTGC TGTTTTAGAA GGGACTGNCA ACTTGCTGGG GGTTAT

SEO ID NO:1558: (Length of Sequence = 292 Nucleotides)

AATTCCCCCT TTCCAAATGT ATTTCAATC CCTTGAGTGT CTAGGCTTCC TGCTTTTAAG GCCTNCCTTC TAACCCAGGG
TTGCCCCCATT CACCTTAAAA CATTTTCAA TAACCCAGAA AAAACCAGGN TGAACATACC CAAGCTCCGG AACCAGCAAA
TMTTGTTCGA ACCCCGCTGA TGACTCCCAG GGGAAGCCAA GAGGACAAAG ACAAGGATGA GGACGAGGAC CCAGGGACCG
MTGGTGAATG GCAACTGCTG TCAACTTCAC TTTTCAACCT CAGNCAGTTT GT

SEO ID NO:1559: (Length of Sequence = 246 Nucleotides)

GENERAL CICAGOCCAA CAAGAGIGAT COTTITAAGG TOCACACACG CIGOCTOTOC TECHLOLICA IGAGOCTOTG GCATGGOCTOT TOCTOCAGCT GGCCCCGGGC TGGGCAGAGG CICCTCCTGC CGGGGCCCCT GCCCACCCCC TCCTTTGCCT GGAGINAGGG TGTTCATACC AAAGACGGAA CCATTTCGCC TTTAAAGAAA ATATATNCAG AAGCAGCCGC TGCCTCGNAG CCCTGG

SEO ID NO:1560: (Length of Sequence = 383 Nucleotides)

CCAAAGGIAC AACAGATTIA CIACATTIAA GACAGGAATC TITICIAATC TCIGIGCCIA TIAAAGAAGC CACCIGCITA GAAGTACTIT GIAGATGAAA AAATACITAT GAATCCACIG TAACITCACA ATCITGAATG CCCAAGGAAAA ACTITACIAG TITICATTIAC CACTATTCIT TAAAGINCIT TITIGATTITA TGITTIAAAAT TITITAATTI TATATTITGA GACAAGGICT TGCICIGITIG CCCAGGCIGC GGGGCAGTGG CATAAACGIG GCICACTGIC ACTITGACCT CCTGGGCTCA AGGAATCCTC CCATCTIAGN CICCIGAGCA AACTGGGNCC ACAGGCATGC ACCATCATGN CCAGCTAATT TIT

SEO ID NO:1561: (Length of Sequence = 313 Nucleotides)

CCCCCTCCAC CGCAGTCTGT GCCCCCGTCC CCACCACCAC CTTCCCCAAC CACTTACAAC TGCCCCAAGT CCCCAACTCC
AAGAGTCTAC GGGACGATTA AGCCTGCGTT CAATCAGAAT TCTGCCCGACA AGGTGTCCCC CGCCACCAGG TCCGACACCG
TGGCCCACCAT GATGAGGGAG AAGGGGATGT ACTTCAGGAG AGAGCTGGAC CGCTACTCCT TGGACTCTGA AGANCTCTAC
AGTCGGAATT NCGGCCCGAA GNCAACTTTC GNAACAAGAG AGGCCAGATG NCAGAAAAACC CATACTCAGA GGT

SEO ID NO:1562: (Length of Sequence = 320 Nucleotides)

ANACGGGCCG CGAACCGCAG TATCATGCTG GCCAAGAAGA TCATCATTAA GGACGGAGGC ACGCCTCAAG GAATAGGTTC TCCTAGTGTC TATCACGCAG TTATCGTCAT CTTTTTGGAG TTTTTTGCTT GGGGACTATT GACAGCACCC ACCTTGGTGG TATTACATGA AACCTTTCCT AAACATACAG TGTGTAACAG TTCTAATACA GCAAATTTAA TACAATTTTT TATTAGATCA AAATTCAATA GAATGTTCA TATGTTTTAA GGAAGGTTCA TTGAATTTCT TCTTTTCAAT GGAAGTCTTC ATTTGGAAAA

SEQ ID NO:1563: (Length of Sequence = 299 Nucleotides)

GCACAAGCAT GACCIGAACC TGICACCIGC CCGINAGIAT TICACATITC TATAGITITI TGIGATICIG CCIGCATITA ATCATCATCA CCAACAAAAA TAGITCCICT GAAGAATIAT TITATACTAG GATTCICAGG NIATCICCIC TCAATCICIA TIGGGATCAC TCCACICIGA CITGIACACI CATITICCCA CIGATGIAGC TGITCICAAG TIAGAAGITA AGITCICAGI CITCATITITA TCAGICATCI CAGCAGCATI CATIATGGIT CAGGCACCICC CICCIATTT

SEO ID NO:1564: (Length of Sequence = 325 Nucleotides)

CAGATGENIC AGITCATACT CIGGCAGITA ATTITATITC CICTAAATAA AAATGGACAG GITAATITAT TAAGCAGCIG
TGITATCAAT AIGGIACGIG TGIGINCITG TATAGATAGA TGIATATGIA CATACATAAC TATACATITI NCIGGACACA
TAATATITNA GGIGCCIATT GIATGCIAGA CACIGITCIA CCATCAGIAA AAAAGCACIG CCCIGITTIA CIGITGATIA
AAAACAAAAT TCIGAAAATA GIGANCAATG AGGCTTACAA CATITGITAC AGGNIAAGGN ATCICAATIT AGGAAAATGI
TGICA

SEO ID NO:1565: (Length of Sequence - 382 Nucleotides)

TITITITITA TATTAGIGCC TGCTTTTAA AAGITTATTI TACATTITAA ATACAGIATI TITCICATAA AAAAAAAATC CAGGAAGIGC CTAACTCCAT GGTTTCTATA CCATATGIAC ATGAAAGCTG ACAGAGAGCC TGACAAATGI TCTGGATGIA ACAGTATGAA CACCIATGAG CTGGGACTAC TTCTGANICA AAATTAAAAA ACACAAATTA AGCACTGCTT AAGAAAAAAA AAATCCAGTT TCTGAACAAC CAAAAGAGAA CAGGGTTAGA TATGTACAAA ACCAGGTATT AAAAANCAGN AAGGAATACA GCACCACAAAA ACTCAAACAN CCCATATGIA GTGAACTGTA TATACTGCAG TTAATGAAAA CC

SEO ID NO:1566: (Length of Sequence = 305 Nucleotides)

GCACTGTGGC TAATTGTAGC TCAAAAGATC TGCAGAGCTC CCAGGGCGGA CAGCAGCCTC GGGTGCAATC CTGGAGCCCC CCAGTGAGGG GTATACCTCA NTTACCATGT GCCAAAGCAT TATACAACTA TGAAGGAAAA GAGCCTGGAG ACCTTAAATT CAGCAAAGGT GACATCATCA TTTINCGAAG ACAAGTGGAT GAAAATTGGT ACCATGGGA AGTCAATGGA ATCCATGGCT TTTTTCCCCA CCAACTTTGT GCAGATTATT AAACCGTTAC CTCAGCCCCC ANCTCAGTGC AAAGC

SEO ID NO:1567: (Length of Sequence = 292 Nucleotides)

GATTICCCTG GGGAAGACAA CATCACCAGC AAATGGATGA TIGTCAACTG GGGAGCCATT GACTCTCCAC TIGATTGTGG
GTTGAGGTTC TNCTTCAGCC TCACATAACA AGATGCCATT GCTTCCGGTG CTATACACAG CACTCTGAGG CTTCTTTGTC
CAGCGAGGAG GCTCTTCTAC TATAACGTGA AAATCGTGAG TGGCTGTTCC CAAGAAATTG CTGGCTGTGC AGCGATAATT
TCCTTTGTCC TGGTAGGAGA CATNCTCTAT CTTCAAAGTC TTGCCATAAT TT

SEO ID NO:1568: (Length of Sequence = 204 Nucleotides)

ACCIACTCAG GAGGCTGAGG CAGGAGAATA GCTTGAACCC AGGAAGCGGA GGTTGCAGTG AGCCGAGGTC ATGCCACTGC ACTCCAGCAT GGGCAATAGA GCGNGACTCI NTCCCCCCGG AAAAAAAAGAA CAAGGGCTAA NTTCAAATCA AATTTTCCCT GTACCCTAAG AANAATAATT AGGNCGGGAG ATGTTTGACT AAGT

SEO ID NO:1569: (Length of Sequence = 362 Nucleotides)

CACAAAGCCA AGTACAGAAC CACAGAATGA AGCCGTCACA AATGTTGAAT CCCAAAACAC TAACAGGAAC AACTCGTATT
TCCATTAATC AAGATTTTAG TATACCAAAT TTTCTAGTTT TTATCTCATG GAAATATAAG GGTATTTTAT CTTTTGTATG
CTACTGAAGG GNAAACATCA TCATACAGCA ATGAATACTT CAAGGGNCTT GTTGATCTCT CTATTATTGA CAGTGGGGTG
TTAAAGTCTC CCACTATTAT TGTGTGGGGG GCTACANCNC TTTGTAGGGC TCTAAGAAGG TGTTTTATGA ATCTGGGGGC
TCCTCTTTGG GNGCATATAT AATTTAGGGT AGTTAGTTCT CC

SEO ID NO:1570: (Length of Sequence = 262 Nucleotides)

TGCTAAATGA TAGANGACAG ATTCAAAGIT GIAGITACTG CGIAACTITA TITATGAGGC ATTITAGAAT AGGCAAAACT GATCINITGT GGTAGAAGTA AGAAGTGGGG TACCCTCTGG AGGAAGAGAA TITNCTITGA AGTGGCATGA GAGGATTITT TTGGCTAATG AAATTATITT NATATCTGAG TAGGGTTGTG GGTTACACAG TITAGGCATT TNTCAAAACT CATGGNACCA TTCATCCAAG TCCTGTGCAT TT

SEO ID NO:1571: (Length of Sequence = 402 Nucleotides)

TGCTAAATGA TAGAAGACAG ATTCAAAGIT GTAGITACTG CGTAACTTTA TITATGAGGC ATTITAGAAT AGGCAAAACT
GATCTGITGT GGTAGAAGTA AGAAGIGGGG TACCCNCTGG AGGAAGAGAA TITNCITTGA AGTGGCATGA GAGGATITGT
TTGGCTAATG AAATTATITT TATATCTGAG TAGGGTTGTG GGTTACACAG TITAGGCATT TGTCAAAAACT CATGGAACCA
TTCATCCAAG TCCTGTGCAT TTTACTGTGT GAAAATTATA TCTCGACTTT TTTCAAAAAA GGAAAAAATA CTTAATTATA
ATATAGCATT TATGNATTAA AATAATCCCN TTATGTAAAAA ATATTTTATT GGNITGGTCA AGATTCATGA TTGCAAACCA
CC

SEO ID NO:1572: (Length of Sequence = 417 Nucleotides)

CTACCAGCCC GTTTCACAA CTAGCAGCAA ATCCTGAAGC ATCCTTAGCC AACCGCAACA GCATGGTGAG CAGAGGCATG
ACAGGAAACA TAGGAGGACA GTTTGGCACT GGAATCAATC CTCAGATGCA GCAGAATGIN TTCCAGTTATC CAGGAGCAGG
AATGGTTCCC CAAGGTGAGG CCAACTTTGC TCCATCTCTA AGCCCTGGGA GCTCCATGGT GCCGATGCCA ATCCCTCCTC
CTCAGAGTTC TCTTCTCCAG CAAACTCCAC CTGCCTCCGG GGTATCAGTC ACCAGACATG AAGGCCTGGC AGCAAGGAGC

GATAGGAAAC AACAATGIGI TCAGTCAAGC TGICCAGAAC CAGNCCACGG CTGCACAGCC AGGNGTATAC AACAACATGA GCATCACCGT TITCCAT

SEO ID NO:1573: (Length of Sequence = 368 Nucleotides)

CAAATAAGTT AGAAACATGA AAAATTCITA GAACTTTAGA TGAAAAATTA AATTTACTAC TAATACCCAC CIGCAATAAT
TTCCCGTAGT TIGGGATCTA GGTTTACAGT GCATGGCAAA AAGACTTTTA CATCTCGAGC CACAAGAACT GGGGTCCTTG
AAGACAAAAA CACTTCAAAA TTTCTTATAT CTCCATCAAT TTCAAGAAGT GGCTCAACAT CCTTAGTTGT TGGAATATTC
TTTGATATTC TTTCGTAGAT GGTTTTTAAT GTCATTTGAT CTGGAATACC TTCAGTCTCT TCCAAATATA ATATGAGNCA
TGAAGTCCGG TATGGCCACT GCTCAGTAAG GTTGATCCCG CTAGCAAG

SEO ID NO:1574: (Length of Sequence = 397 Nucleotides)

AATTTAAGC AAATGTATG TITAAAGACT GTITTGATGA AAACTTITAG AATTGAGTTA GTAGCAGAAT ACATAGCTAA
ATGTACTTIN CIACAAATAG AATGAGATAT TIGATTAAA ATATINCTIT CCTCTTGAAA TAGGATGTTA GATAGGGACA
TCTCATTITA CCTATCAAGT TCTGAGTCTT GCTTTAGAAC TACTTCTTTT AACTTAATTN CATGCATACA CTGGAGACA
ATAATATGGC TTTTTAACTG CATTATCTTT AGTTGAAACT GATGGAGAAA CAAAAATACT GCTTATACCA TATTGGGTAC
ATGCTGAATG TTTTTAAAGA CTGCCAAAA CTGACATTTT TTAAAATTAA ATAAGATGTT TTAGTTTCAA ATTAGAG

SEO ID NO:1575: (Length of Sequence = 296 Nucleotides)

GEACTCAGCC TTCCGCGGCA TCTGCATGAT GATCGGTGTC AACCCGGGGG GCGTTGTGCA GGTTGGGGCA GCTGGGCTCT
NAGGGCAGGC GCGGGCNCTG GGCTCGGGCG GCCCCTCACC TGGGATCCGT CACGTTTCAG GACTTTATTT TCTTCTTCAA
TGNTGTAGCC TCCTGGGTGA GCCCGAAGAT NACCTTCGGG ACATGTTTTA TAAGGTGAGG CTCTGTCTGG GCCCTGATCT
AGTTCCGGGA GCAGGCAGGA NGTGAGACCA TCTGGTAACA ATNGGGGCTN GGGATT

SEO ID NO:1576: (Length of Sequence = 289 Nucleotides)

CTITIATGAAG TAGTAATICC TGAGAGGIGI GCTGGCTGAA AACATAATAG GTTCTGGAAG AGCCAGGIAA ATGCCTGGNI TTAGACATGC AGGGGITAAT CAAAATAATI TAGGAGCGIT TTCAGCTGGI GAGCCTCATA TGGGATCTIC GAACCCGTGG CGAGAAGAAA ACCGGIGITT AGGNAGCACC AGGCACAGTG CTCGGAAGGG AGAGGCTNGC CGGCCAGTGI GCAGCTCAGC TMTTTCGAGG ACGGAACCCG CAGCCTNGCT GTNTCCCAGC AGACCCAGG

SEO ID NO:1577: (Length of Sequence = 320 Nucleotides)

CAGACTCTAC TCAGATTTCC CGCCTATGCC CCTAGGACAG AGCTGGAAGG GAAGGAGGCT GGGCCTATTT AGTCATAATG
CCTCCCCACC AGGTCTAGCT TTCATTCATC CATGAACCCT CACCCAAGGG CCAAGAACTG AGTTCACTGC ACCCTGGACC
CCTGTTGAGG TAGGAGAAGT AGACGTTGGG AGCAAGGTTC CTCTCCTAAT TTTNTTGCAT CCCCTCAGTG CCCAGCACAG
CTCCCGGATAC AGGGCAGGTT CACAGTCAGC GTGTTCACCT GGGNCTGTGT ATGCACCTAA GGAAAAGNCT CAATTTTCCT

SEO ID NO:1578: (Length of Sequence = 217 Nucleotides)

AATCAGGAGA ACTGTTAGAG CCATACCAGA GAAAATCACA AGAAAGGCAG GACTGCAAAG NTCTAGTGGA GGCTGTGAGA AAAGGTAAAC CCCTTCTTAA GCTCATCTGC CCCTTTAGTT ACCACTGGCT GTCTCACTCC TGGATTTATG TGACTCCCTT AGCTATACTT TCCCANCCCC CTGGGATGTT CCCCACTCAT CCCVATTCACT CACAAAG

SEQ ID NO:1579: (Length of Sequence = 375 Nucleotides)

TIGGICCICA AGICCIATIT TAAAATTIIG TCAATTAGAG GACICTIGGI TCICTIGGIT GACICATICT CIGCIGATIT
GITCICIGIA CIIGCAGCAA ATAAAGIGCA GICATIGAGA ATGINCCIGI GICACIGIGA TGIATCAAGG GATCITCATG
TIAATATCIG TITCICIGAC AACTGIGITT TATACITIGI ACTGIAGCTI TCATIGGAGA AGCCCIGGGC TCATAAGAGT
GATTIGITGI GGCATITCCT TATGGAACAT AAGCTITIGA AATATACTIG AGGTAAATAT TCATGGGAGA CATCCAAATG
CAGIAATGAG AGIACAATGA AGACAGCATI TINGACTITIG GAAACCIGAG TICAA

SEO ID NO:1580: (Length of Sequence = 325 Nucleotides)

TCINCIGATG CACCCATGAG AGGGGAGACA GCACTGICGI CICTCGCAGI TITCCCITAA CACTCCCITA TCTGCAGACT
TAAACTAGGA GCCCCIGGCA GAGICCIACC TCCAGAATCA CAAAAGIGTA GAAGGAAAGI GAGAGACATT GATTGACTIT
ATATCTGACT TACTAGITIC CIAAGGCAGA GATTTITIAG AAAACTGCCI GGCCIGGCCC AGCCCAGGAT AGATAGGGAT
GGGTAAGAAG CCCTINAGAA TGTGGCAGIA TGTGGCTING ACTTCAGACT TGTCAGATTA GGGGTTTITT
TTAGC

SEO ID NO:1581: (Length of Sequence = 402 Nucleotides)

GCAGATCAAG AAAAAGITIC AGCCAATGAA CAAGATCGAG AGGAGCATAC TACATGATGI GGIGGAAGIG GCTGGCCTGA
CATCCITCIC CITTGGGGAA GATGATGACT GICGCTATGI CATGNICITC AAAAAGGAGI TIGCACCCTC AGATGAAGAG
CTAGACTCIT ACCGICGIGG AGAGGAATGG GACCCCCAGA AGGCTGAGGA GAAGCGGAAG TIGAAAAGGAG CTGGCCCAGA
GGCAAGAGGA GGAGGCAGCC CAGCAGGGC CIGIGGIGGI GAGCCCTGCC AGCGACTACA AGGACAAGIN CAGCCACCTC
ATCGGCAAGG GAGCAGCCAA AGACGGAGNC CACATTCTAC AAGGCCAATA AAGACCTACG GCTTTITTCC CNIGGCCAAT
AA

SEO ID NO:1582: (Length of Sequence = 286 Nucleotides)

TCTTAGTIGA TTAAAACAAA TAATIGAAAT AAAAAATTAT GITTATNCIT ACATGIATGC CATGIAGCAC TITAAGGAGA TGAGTTATG AAATTCATGA ATGAGAGGAT GATGIAAGIT TAAAAAATCAT TATTTTAGTI GCTTTATTCT NCTATTITTAA ATTCATAAAT AACACAGGIG GCCTGIATTI TGAAAAGAGC CCTTTCCTCC ATTTGANCTT TATAAACACT GAGGCAGTAG GIGIAAAATA TTATCTCCAC TTTATATTIG AAGGAAATGG GGGCCA

SEO ID NO:1583: (Length of Sequence = 323 Nucleotides)

CTAATTTTG TATTTTAGT AGAGATGGGG TTTCACCATG TTGGCCAGAC TGGTCTCAAA CTCCTGACCT CAGGTGATCC GCCTGCCTTG GCCTCCCAAA GTGCCAGGAT TATAGGCATG AGCCACCACG CCTGGCCTTC CAGTTGTGAC CTTGTTAGGA TACTGCTTTA ATTCATTTTC CCATTGAAAA TAAGCATGAA AATAACTGTG CAGTCATAAT TGTGGTATTT NCTGTNAAGG AAAGTGGCAG GGCTCTGAGT GTTTATCGGG AGACCTAACC CAGTNTCAGA GGGGAAGTCA GAAGGCTTAC TNCCCAATGG GGG

SEQ ID NO:1584: (Length of Sequence = 301 Nucleotides)

AAATACTIGI AAATCACTIT ATGTTTCIGA GTAAGGAAGI AATGAAACAT ACGTACAAGI AATCAGTAAG ACTTGITAGA CAGCTGITGT TCAGGATGCC TITTAAAAGGG CIGGTAATGC AGTTACATIC TAACAGAGAA GTCCAAACTA CAGGTAAAAA CTACGGCTIG TACIGIGAAA AATGTGCAGC TITTCAGTTA TAAAACTAGT TGAACACTGG TTTACAAGGT AATCCGTAGG AACAGAGAGA

SEO ID NO:1585: (Length of Sequence = 328 Nucleotides)

AAATACTGAT TTCAGACCTT CTTGCTCTAG AAGTCAAAAT ACTTTCCCCC TGACAAGAGG TAAGATAAGG TAGAAAATAG
AAACACTGGA AGAGAGATCT GGACTCCTAA AGCTGTGATG CCATAGTGTA GTGGGGGGGG GTGCGTGAGG AAGTCAGGAA
TGCCGCCAATG TTAAAGGGAA AGGGAAGATG GAGCAAAGTG AGTCCCAGGG CCAGCAGGGG GCCAGCCTTN TTTGACAGGG
GCAGGGGAGA AAAGGCCAGA CTTCCCATAC ACATGCTAGA GGGGAGGGCT AGTGTTGAAG GGTAATAAGT TGAAGGAGTC
CACGGGCT

SEO ID NO:1586: (Length of Sequence = 256 Nucleotides)

GGACTATCTG TATGGCAGAC TCATCAACTT TGAGAAGAGG AGGAAGGAGT TCGAGGTGAT CGCCCAGATC AAGCTGCTGC
AGTCGGCCTG CAACAACTAC AGCATTGCGC CAGATGAGCA ATTTGGGGCC TGGTTCCGGG CCGTGGAGCG CTCAGCGAGA
CTNAGAGCTA CAACCTGTCG TGCGAGCTGG AGCCCCCATC CGAGTCAGCC AGCAACACCC TCAGGACCAA GAAGAACACA
GCCATINICA AGCGCT

SEO ID NO:1587: (Length of Sequence = 371 Mucleotides)

GGATICIACA GGCATAGACT TACACGAGIT TCTGATTAAC ACATTAAAGA ATAATICCAG GGACAGGATG ATACITITGA
AAATGGAGCA GGAAATTATT GATITCATTG CTGACAACAA TAATCATTAT AAAAAGITCC CTCAGATGTC ATCGTATCAG
AGGATGCITG TCCATCGAGT GGCAGCTTAT TTTGGATTGG TTCACAATGT GGATCAAACA GGNAAATCTG TTATCATCAA
CAAGNCCAGC AGCACCAGAA TNITACCAGC CAGTCTTGTC TNGTCAACAG GGGNTTCCAA GGGCTAATAG GAGTNCAGCA
GCCCACCTCA GAGTCAGACG TGGTTAAATN ACCCCCAAGG GACTCCGGTG C

SEO ID NO:1588: (Length of Sequence = 314 Nucleotides)

CACACAGGAT TOCATAATAC TOCTGCTGTG TTCTGAATAT TTGTACTTCA CATGGGATTA CTGAACACTA CTACGAGATT
CTGAATGTTT GINGCTCACA TAGGATTCCA AAATGCCCCT GCTGTGTTCT GTTTGTCCCT CACATAGGGT CACTGCTGCT
GGGTTCTCAG TGTTTCTCAC TCACATAGAA TTCCAGAACA CTGCGAAGAA TTTCTGAATG GTTTTCTGTA ACATAGTATT
CCAGCACACT CTCGCTGTTG TTTGAATGTT TGTCCCTCAC ATAGGATTCC AGAACACTTC TGCTGATGTC TTGA

SEO ID NO:1589: (Length of Sequence = 256 Nucleotides)

GACGAGGCAC CATGCGTGAN ATCGTGCACA TCCAGGNGGG CCANINCGGC AACCAGATCG GNGCCAAGIT TTGGGAGGTC ATCAGTGATG AGCATGGGAT TGACCCCACT GGCAGTTACC ATGGAGACAG TGATTTGCAG CINGAGAGAN TCAATGTTTA CTACAATGAA GCCACTGGTA ACAAATATGT TCCTCGGGCC ATCCTCGTGG ATCTGGAGCC AGGCACGATG GATTCNGTTA GGTCINGACC ATTCGG

SEO ID NO:1590: (Length of Sequence = 313 Nucleotides)

GGCAACAAGC CAAGTAGCAA AGATATAAGC AACAATCAAA TGGAGCCTGA AATATGATAA GAGCATACAT GCACTITAAC
AATAATTIG ATACTGGAAT GATTATTICA GAAGCAATAT TITINCIGAA AAGCATTGGT CTICTGTACA GAAAAATAAA
AAAGTGAGCT GCCACTCATA GIGAATTAAG AGCTGTGGGC TGAAAGGGTC TCTTTTATAG CCAGTTTGAA ATTTTTCATA
TAATAAAAAC AGTATGTAAA TATTATATAT ATATACACAC ATACATATAT ATGCATATAT GTACATATTT CTG

SEQ ID NO:1591: (Length of Sequence = 296 Nucleotides)

TTINAGICIC CUGCCICACA ATTCAGCGAC TGCAGCICGG CCAAGGCCAG GGGAGACCIG GGIGCCITCA GCAAAGGICA
GAIGCAGAAG CCATATGAAG ACCCCTGGIT TGCCCGGCGG ACMGUGAGAAAA TGGCGGGAGACCIG GGGGIAGGAIT NGGGGCCCAG GCCTGGCCTC GGGGTTCCCC CGCTGCCTGC TGGCCAGTGG
CNGAACCCCC CANINCCTGC CACINTCACA CAGTATITAT TGTTACCAAA ATGGCT

SEQ ID NO:1592: (Length of Sequence = 299 Nucleotides)

GGAATTCCCA AATTATGGGT AGTCCAAAAG CCAAAGGCAA TGTGAGGAAG GACACTCCCC AGATAAGAAC AAAAACAGAA ATCTGTATGT NCTATGTGTT ACACACAGIT GCGAATAATC AGATGTACAC ACATGATGCA AAGGCACGCC GCTACACATT TATGTGATAT TCAGACATAT GTTCAAATAG AGGAGGTGAA TATCTTTTTA TAAATACAAT TTAGCAAGTA CAAGAATGCT GATCAGCTGC AGCTCAAGAG GAAAGGGGGG AAAAAATCTT ATGGGAAATT ATTAATACT

SEO ID NO:1593: (Length of Sequence = 378 Nucleotides)

CCAGITTGGT GATTCINITC TGIGTCTGCT GATCIATTGG CGIGAGAAGC TGAAAGIGAC CAGCCAACAG CCATAACTIT
ATGITTAGIG AGACTCATAA TGGGTCTCCT GCTGGAAGAT CTCCCCTCTA AGANTCAGTA ATTCIAGACC TGCAAAGITT
GAAGITGIAA GCATGGGAAA CACAAATTCC CCAAATAGGT CCAGATAGIG ATAGAGAATA AGACACITAC TTGCCTACTT
CCATTCCTCA GCCCAGATAT TCTACCTATA GTGGACATGC CCATGCAATG GGCTATTGGG TTTGAGGTAT ACATTGCACG
GTTGAAGGAC AGTGCCTCAT CCTTGCAGGG GTGCCCTTIN CCAGTTGGCA CCACAGCT

SEO ID NO:1594: (Length of Sequence = 353 Nucleotides)

ATTITINCEG GGGAGGIGIA TGTAGATGAG AGICTATGAT ATAAAGCAGT AAAAAAAATG CIGITGTATA GGGATGCAAT
ATTITCGGIG TAAGGAAGAG GITTTAATTC ATAAAATAGA AAACAGGITG GAGAAGTCTT TAGGAAAGGG ATACCITITG
GGTTGGCTTT TGAAGGAGAA GITTATACCC AGGITCAAGC TGAAGGGCTA AGIGAGTAAC TGAAAGGGCT GAGCTATTIG
GATTACCATG AGGAATTIGT GATGGCTGGG AATGTAGGGT GTGTGACCAG ATGTGGAATC ACAGAGGGAG CCCACAGAGG
AGCTTCGGCA CATAANCTAA AGAGTTTAAT TIT

SEO ID NO:1595: (Length of Sequence = 343 Nucleotides)

CAATATATIA AATCIATTIT GIAGCIGGAC TICACITACA AIGIAACAGA ACATIGAATA TIAGATICIG AGCATATICA
TGCAAACITC CACITIGGIG AAAGIGAIGA CAGIGGAGIT CIGGAAGACA AITITICCTIG TAAACACCAA GITTIGCACI
TIGGACIAIG CICTCAAGAT AGAAACITAC GIGAGIGGAA AAAGAAAATG TATAAATGIG AACAAATATT CCITACCACA
CAGAATAACC CIGGCAACAA ACAATATCCC CAAGICCIGG GINATICAAT CCICACCGIG GGCAGGAAGG GIGAAGGAGG
CTGCACCIGG GNCACAGCCT TIT

SEO ID NO:1596: (Length of Sequence = 373 Nucleotides)

TAGICAGITA TIGCIGCACT AGAGCTAAAT AAAAGACATA AATATCTAAG GCACTTACTG GAATAAACAT CITATITICCG CIAAGAGGIT GGCTAGGGAA GCTCTGCTC AGAGTATGGG TIGAGTATAA GCCTGTNCCA CATGTCTTTT GCTCTGGGAC CAGGAGTTGT GCAGCCCATC CITTTCTCAA GACAAAAGCT GAGCCAAGCA AGGACATTTA AAGCTTCACT TCTGCTCACA TCATATCTAT TGGNCAAACA TTCCATTGGG CCAAAGCAAA TCACATGGGC CAAGTCAAGC ATCAGTAGGT CTGGGGGGAAT ATTCTTTCCT CTACTCTTGG ACACATGGGA AAGGGTTATG CATACTAATT CIT

SEQ ID NO:1597: (Length of Sequence = 276 Nucleotides)

GATTGICCAT ACITGATTAT TAGITTCTAA AGAAAGTATI CITAATTCCA AGCCTAATAG CICITATGIC ATTAGITTCT AGTGCAGAGA AATGTACTTG ATGAATTTTT GITGACTTTT TTTTTTTGCTA GCCAATATGA AGGTTGCCAG TCCCTGCCAA LACACACTA AAACTATTT TNCATGAGTA ATAACAATAA TATTCITTTT TAAA1AGCAC CITTAACCCA AAAATCTTAA GCCTATATAA ACATTCACTC AACANTACAC TCAAAA

SEQ ID NO:1598: (Length of Sequence = 355 Nucleotides)

TGIATIGCIA ACTGICITIG TAACIAATIT ATGIATACNC TAAATGGIAT AGCATGIGAT TITATIATAG TIGATIAACI TIGIAATINC TGIAACIGCA TOGATATCCC AGICTACCTG GAAAATIAAG TCTATIAACC ATAGITGCTG TGGGAGACAG TACTATIGCC AACTGAAGCC TGAATCCTTC ATTTATITTG TCCCCAGITA CAGAGIGGAG GITTAGAGGA GIGGGGITAG ATAATGCTCA GATTAGAAAT ACAAAGGCAG CIGICAGATC CTCCCATITT ATTTGITTGA AGGAACIGAG GITGGTAAAC ATCACAAGNG CIAGITAACT GGIGAGIAGC AGCCC

SEO ID NO:1599: (Length of Sequence = 313 Nucleotides)

GGAGGTGAAG GACACAGTGG ATGGGCAGAG GNTCCTGGAG AAGAAGGGCA GTNCTGCNCT CAAGGACCTC AAGCGGCANT GCATTTGGAG CGGAAACGGG CAGATAAGCT GCAGGAGCGA CTNCAGGACA TCCTCACTAA CAGCAAGAGC CGCTCAGGCC TINAGGAGCT GGTTCTCTCA GAGATGAACT CACCAAGCCG GACCCAGACA GGGGACAGCA GTAGCATCTC CTCCTTCAGC TACCGGGAGA TCTTTCGGGA AAAGGAGGAG CTTCGGCTTG TTCCAGCCAG GTCCTTATCC AGCAGNCCTN AAG

SEO ID NO:1600: (Length of Sequence = 277 Nucleotides)

AGTICACAGA ACTOCAATIC TITATTAATO ACAGCITECT CACAATGACA TACAGGAAAA TAGCACTAAT GAAGAGTAAA
TATGCAGGCA GCAACCITCA GGAGTIGGGA GITGGGGAGA AACGACTICA AAACTGCGAT AGGTACTTAT GGTGGGTATC
TGGTGATTCT TAGTTGGCAC AAATGCCCTG CCTAGCCCCC TTAACTGCGT CANITTCACA GATGGAGTGT TITGTTGTTG
GTGTTGTTAG TAGGCAGGAT TGCCTTACAC TGGGGGA

SEO ID NO:1601: (Length of Sequence = 228 Nucleotides)

TTGAGACCAT CCAGGCTAAC ACGGTGAAAC CCCGTCTCTA CTAAAAATCC AAAAAAAAA AAAAAAATT AGCCGGGCGT GGTGGCTTGC GCCTGAAGTC CCAGCCACTA AGGAGGCTGA GGCAGGAGAA TGGCATGAAC CTGGGAGGCG GAGTTGCAGT GAGCCGAGAT CGCGCCACTG CACTCCAGCC TCGGCGACAA AGCAAGACTC TGTCTCAAAA AAAAAAA

SEO ID NO:1602: (Length of Sequence = 299 Mucleotides)

GGAAGTCCTT TCTAATGAAG AGGGGAGATG TTATCGATTA TNCATCATCA GGGGTTTCCA CCAACGATGC TTCCCCCCTG
GTTCCTATCA CTGAAGAAGA TGAAAAATCA GATCAGTCAG GCAGTAAGCT TCTCCCCAGGC AAGAAATCTT CCGAAAGGTC
AAGCCTCTTC CAGACAGATT TGAAGCTTAA GGGAAGTGGG CTGCGCTATC AAAAACTCCC AAGTGACGAG GATGAATCTG
GCACAGAAGA ATCAGATAAC ACTCCACTGC TCAAAGGATG ACAAAGACAG NAAAGCCGA

SEO ID NO:1603: (Length of Sequence = 263 Nucleotides)

AAGGCAAGAA ATTAGCCTTG TTAAGAATTT TAAGTGTAAT GGGAAGCCAT TAGAGGGFTT TAAACAAGGA AAGATGTGAT GTGACTTATA TTCTAATAGG ATTGCCTTGA TTCACCTATG GAGAATGGAT INNTGGGATC TCAGTACTGG GATACTGAGA TCCCCAGGGGG AAAATATCAC TAAGGTTGGA ATTGCTTTTC TGCACATTAA AAGCAATTCN CTTTTTCCTT GAAACCTCCA TGTGATGTTA ATTAGGGTAA ATG

SEO ID NO:1604: (Length of Sequence = 260 Nucleotides)

ATGAAGACGT ACGACTTATT TTTGTGTTCT GAACATAAGT NCTTTGTCAC ATAAAATGTG CTATGAATGT TGAGTTTTAA
ATACTCGAGC GGTGACTCAC GCCTGTAATC CCAGCACTTC GGGAGGCCAA GGCGGGCGGT TCACCTGAGG TCAGGAGTTC
GAAACCAGTC TGGCAAACAT GGTGAAAACC CCGTCTCTAC TAAAAATACA AAAGTAGCGG GGTGTCGTGG CGTATGCTGG
TAATCCTAGG GTTCCTGTCA

SEO ID NO:1605: (Length of Sequence = 290 Nucleotides)

GACAGACATT CAAACCATGG CAGGTGGCAA GAAGTATCAA ACTACTAGAT CCITGGGATT GINCTITGTA CTGGGGTGTA
TTTTTNCCAA CAATCCTAAA AATCATATGA ATAGAGATAG CAATATATAT CTNACCCATT TGGAAATGCA CAGAGATTCA
GGAGTGTCA CATAGAAACA GAAGATCATT GGCTTTTGTC CATTCCCAAC GCCAGNAATC TGTTTTCCTT GACTCTTTTT
GATCTGTGTT TCTGAATGIN TTGATATACT GCGCCTACTG GGTGTGCAGG

SEO ID NO:1606: (Length of Sequence = 290 Nucleotides)

CTCACTIGGG TACTACAGIG TGGAAGCIGA GIGCATATGG TATATTINAT TCATTITITGI AAAGCGITCT GITTIGIGIT
TACTAATIGG GATGICATAG TACTIGGCIG CCGGGITIGI TIGITITIGG GGAAATITIG AAAAGIGGAG TIGATATTAA
AAATAAATGI GIATGIGIGI ACATATATAT ACACACACAT ACACATATAT TATGCATGIG GIGAAAAGAA TIGGCIAGAT
AGGGGATTIT CCIGAACACT GCAAAAATAG AACGTAGCAA AATGCCTICA

SEO ID NO:1607: (Length of Sequence = 365 Nucleotides)

GCTCCACTGA CCAGCTGITC CCTGTCTCC CTTCTCCTTG AGCCTCCTC TTCCCTGAGA CACAATAATA TTAAAATTTG
GCCAATCAAT AACTCAACAA TGGTGTCTAA TAATTGITCA GGTGCGAGGA AGAGGCATAC ATCTCTCACT TTAAATCAAA
AGCTAGAAAT GATTAAGCTT AGTGAGGAAG GCATGTCAAA AGCCGAGACA GACCAAAAGC TAGGCCTTTT GTGCCAGTTA
GCTAAGATGT GACTATAAAG AAAAGCTGTC GAGGGAAATT TAGAATGGTA CTCCAGGGGA ACACACAATG ATAAGGAAGC
AAACAGCCTT ACTACTNGGA TATGGGGAAA AGTTTTCAGC TTTGG

SEO ID NO:1608: (Length of Sequence = 294 Nucleotides)

CTCAGGAAGC CTCTCTTCT TCACTTACCA TTACTAACTC TCCAAGCATA GAAATCCCTG GGAATTGCGA GAATAACTCC CACTATTTTA AAATTTATAT TCAGATTTGT TTCGTTTCAT AAGACACATC AAACAGGCCT ATACAAAAGG TTTAGGAAAA GAAAACAATG GTGAGTCCCG GCCCTCTTCG AATTCACTGG CACCTCATGC AAGINTAGGA AGGCACGCTG GATCGTCTAT CTGATTCCAA AGCTGTCCTT TGCCATCTCA TCCCTTGGNC TGCCCCCCAA CCCT

SEQ ID NO:1609: (Length of Sequence = 393 Nucleotides)

CAAAAGCTAA CTCTTAATAA GAAGATGAGG AAATAAAATC AGTTCAAAAG GGAGGAATAT GCATTCCCAG AATTAAAGGA CCCCGGGTCC AGTTTGAGGA GGACTCTTGG CCAGATACAA GCCCCTTGTA TAATNCTCAA GAGGGAGGAG ACCTTATTTN CTCCTTNGAG GTGTCTAGTA TGAAANCTGC TTATTTTGAA ATGTGATTCT AGCCATTATC AGGRCAACT GCAGATAATT CCCATTTACA GAGGAATGCT GCTAACAGGT GTGGGRGGA GCAGCGACAN CCNAAAAATTC TGCTGTCATA GGTCACGTTT ATGTTGGTTT TCTTTGAAAA TCAAGGGGTA GAAAAATTCA TGCCTCTAGA GGAGAGAGAG GAAACACATG AGG

SEO ID NO:1610: (Length of Sequence = 464 Nucleotides)

TGICIGIATI TATTAAATIG CCTTTACTAC TITTAGATGG CCATACGITI TCAAAAGCAA AGACCTAGIA AGCCATITGT
GITCATITGC TAAGCTATCI TAGGIACAGG TCCAGATTAT AAATGITACC TGCTAATCAG AGAGCAAATI TITAAATTAA
TCACTIGIAA ATCCACATTA AAAGAAAAAG AAACTTAGAA AAACACATAA ATTTCTTITG TGATCCCACT ATTCAGGAAA
ATCCATTGAA AAAGCAGATG ACTTATCCGT GITAAATTIT TAAAGNCCCT ATTTAAACTG TCATGTAAAT TCTNATTTAT
CTAATTTTT AAAACACATA TAGNNITTTA CTCTCCAGTT CCATAANIGN CTCANTTCTG GIGANGGICA TTACAACAGN
CATTACGNGG GCATATCGGN NIAAAANGGC CNIGCGGICC TGNATCNGAG GNGGGGTTAA GGTC

SEC ID NO:1611: (Length of Sequence = 465 Nucleotides)

ATAATTTAAA GAAAAGAGAA TTCTACAATG TAAAACCCTT TAATATAAGC TGTTTTAATA ATTGGAAAAC AGAATGANTA NTGTTTTNT TTGTCATGCC CAATTATTTC ANCAAGTTTT TATTAATAAC TTGCTACATG GTAGGCACAG CTGTAGGTGT TEGAGATATA GAGGIAAACA AGTCTGACAT GATCTATGCT ACCACGGAGT TCTTATTTTC AAAGTGGAAG GTAGAAAATA
AATAAAAATG ANCTAGAAGA GCAAAGTGCC TCTGAATGAG CATGCAGANG CATGTTTTCA AAATGTCTGT GNGTGGGATA
AATAGATCAG CAACACACCA GGCCATGCAA TTINGCAGCA AATCACTTCT GCAGTCTAGC TGCTGTTTTT CCTACTCTGG
AATCATACTC CCCCCTTCGG TCATCINTGC CAGTTTCNCT GNGCTTCACC CTACCCTCCN TTTTN

SEO ID NO:1612: (Length of Sequence = 458 Mucleotides)

ATGAAATTGA ACAAACCTAA AGAGAAATGT TCTTACCGTT CCACAGGAAC CAGCTTCTTC CACTGGGCCA CTAGGTCCCT
GGCAAAGCTT CCAACATGCT CGTGTTTTCG CAAGCTATTT ACTGTTTTCC CAACCCCAGT CTCCTAAAAT TTGACAAAGT
AATTGTTAGA GGGGTCTGGA ACTAGGCTAA CGTTTTTCTA AAGAAATAAG GCTTTCTACT TTGAGAAACT CAACAAGCAA
TACTTCCTTC CTACAACATA CCCTGCAAAT CTTAACACTA AATTACTTTG TGTCTATGNC CCAAATCTCT AATGACACAC
AGTAGCAAAG NGTACCAAGT TCAGAACTTT AATAACACNG GTNATTAGGG CAGGTGTTAG GGCACTAGNT AAGNGCTTTG
CATCAGTTCT GGATCAGNCT TTTAAATAAC CCCTTAAGNG GGGNTNAGNC CCTTTTTT

SEO ID NO:1613: (Length of Sequence = 322 Nucleotides)

ATGIGGAGAT TIGITGIGG CTAGGGCAGI CCAGAGGAGA GATATGIGGC AGGACAAGIC TCIACCCIAT ACAAGINCIT CCGGCAAGCC CTCAGCACAT GACATAGGCC CAGAGAAGGA TGCAAAGAAT TCIGGICATA AATTGITITC AAATATCAAA TAAATCATAT GIGCACATGC ACAAACATGC CITCACAACT GAGIAAAACC AGACTCACCT TCAAATATAT CAACAGTTTT NTCAAGCGCC GITAAAAATC AGGCATCGGA CCICIGGNIN CGAGAGCTGG TTINATGGGG AAGTTAGATC AACCCGTCAT CT

SEO ID NO:1614: (Length of Sequence = 280 Nucleotides)

AGIATCAAGG GATAAAATAT ATTITTAATT TIGIATITCA CITGAAAATT GIAAGGNCCA TITTATAATG TATIGCITGC AAAATAAGTC AIGGAAGCCC TGAAAAATTA GICAATTCAC TAATCAAAGA AACATATATT AAAGACCTAC TATGCAIGAG GCACCAIGCT AATIGCITIG AAGAAGACAA AGITGAATTA GACAGGGNTC CCGITTACAA GNIATITACA AIGCAAAGGG GGATACAAGA CATATAAAAG GCTATGGAAC TGCCCITCCG

SEO ID NO:1615: (Length of Sequence = 393 Nucleotides)

GOGTGGTGGT GOGTGCCTGT AAATCCCAGC TACTACGGAG TCTGAGGCAG GAAAATCCCT TGAACCAGGG AGTCGGAGGT
TGCAGTGAGC CGAGAGCACG CCACTNCACT CCCGCCTAGC GACAGANIGA GACTCCGTCT CAAAACAAAA CAAAACAAAA
CAAAAAAACCA AAAACACTGG GAGTCCCAGT TTGTAGGAAA TCATTAAGAT TTTATTATTT GAGCTCCAGA ACGAGTGAGG
ATGACCTGAT AATTTTGGTT TGGCTCAGGT TGTAATGTGT TTCTGTTTTG CTCGATGACT ACTAGAACAG TTCTCAAACT
GTGTGGTGGG TAAGAATCAC CTGGGGACTT TGACCAAGIN ACATGTCTAC AACACCCGGC CCCTACAGGC TCT

SEO ID NO:1616: (Length of Sequence = 353 Nucleotides)

CCACCCCAGC CICCITGGAG CIATCCCITT CIATCCCCCI CCATCCAGCC CCTGGCCACC ACCATTATAT CIATTCTGGA
ATTCCCACAG GAAAAGCAGG CACTTATAA ATCAGCGAGG GATTCACGGC GAAATGAGAC TGITCGTGAG TNATGGCGIN
CCGGGITGCT TGCCGGIGCT GGCCGCCGNC GGGAGAGCCC GGGGCAGAGC AGAGGIGCTC ATCAGCACTG TAGGCCCGGA
AGATTGINIG GINCCGITCC TGACCCGGNC TAAGGICCCT GTCTTGCAGC TGGATAGCGG CANCIANCIN TTCTCCACTA
GTGCCAATCTG CCGATACTTT TTTTTTGTIA TCT

SEO ID NO:1617: (Length of Sequence = 227 Nucleotides)

TTICITICCAT GCAACANICI GNAGACITAA GIGGCITICI NCIGIACINC CATAGAACCC ACCCAGIACA TACCICCAGI GNGGCACIGA TITTATGCIA TACATATGAC IGIGIGITCA TCTCCICCAC CAGACIGIGA GICCCATIGG AGTAGGAACT AAATTIINIT CAACACTCIG TCTTCATCAC CICGIGIAGI ATCTTGTACA GAGTAGATAA TGATTAA

SEO ID NO:1618: (Length of Sequence = 362 Nucleotides)

GGAAGGTTTT TAATGCATGA NGTATACTTG TNATCCTGGA GGTTGGAAAA GATTCAGTAA AGATAAAGTT TGGCAAAAAT
GATTCTCTCC CTAGGATTTG GGGATATGTA AATCAAACCA AAGGCACATT CTGCAGCTCA CAGCAACCTT CATTTTTTGT
CCTAGATTGA GTTATCTATC AAGAATCATT CATTCCCTCT CAGCCCTTGC AACTGTTTCC TATGACTTTG GACTTGGCCA
TGCAACTTGC TTTGGCCAAT ACAATGTGAG TTAATGTGCT TTAAGTGCAT GTAATTAGGT CAGTCCCTCC CTCCTTGAGC
TTCAACTCTC CACCATGAGG ACAACATTGC CCTCCTTCCT GG

SEO ID NO:1619: (Length of Sequence = 344 Nucleotides)

GCAACCTCAT CCCAGGITCA AGINATTCTC CTGCCTCANC CTCCTGAGTA GCTGGGATTA CTGGCGCACC ACCACACCCG
GCTAATTTTG TATTTTTAGT AGAGACAGGG TTTCGCCATG TTGGCCAGGC TGGTCTTGAA CTCCTGACCT CAGGTGATCC
ACCCACCTCA GCCTTCCAAA GTGCTGGGAT TCCAGGCATG AGCTACTGIN TCGGCCCAAA TCTTTCTTAA GTTGTGTCTG
GCCTTTGGCA GAAATAGCCA CAAAGNCAGG GTAGGAACGT TTTACTCTTC AAGTGATGAT GGCATCCGAT AANCTTTTAG
AGGGAGGTTT TTAAAATGCA ACGT

SEO ID NO:1620: (Length of Sequence = 379 Nucleotides)

GCCAGCOGAA GCTCCTCAGG CTCCCACCCT CTACAAGCTC CTTCTGCTCC AGCCACACTC ACCAGGCCCG AGTTCCCACC
TAGCACCTTC CCTGGGAATN ATCTCCCCCT GGTTGGCTCT TTCTACTTAT TCAGCCTCAA ATGINATCTC CACTGANAGG
CCTTTCCTGA CCTGCTGAGC TTGATTCCCT CCCCTCCCCA GINACATTAC TCCGTGTTAT GGTACCCATC CCTGTCTCCT
TAGCTTGTTT TTGTCTGTAT TGGCTCTTCC ACTAGACTGT AAGCTGCATG AGGGCAGGGG ATGTCTGTTT AATNCCAGTT
GCTCAGGATA GTGTATGGCT CGTGATAGAT GCCTAGNACA TTTTAAAATG GGGACGGAT

SEO ID NO:1621: (Length of Sequence = 283 Nucleotides)

GATTIGGGG CICGGGGAGG CAGAGAATCI CITGGGAGIC TIGGGIGGGG CIGGIGCATI CIGITICCIC TIGATCICAA AGGACAATGI GGATTINGGG ACCAAAGGIC AGGGACACAI CCCCTIAGAG GACCIGAGII INGGAGAGIG GIGAGIGGAA GGGAGGAGCA GCCAGAGAGCA GCCIGITITC ACTCAGCTIA ATTCICCITC CCAGATAAGG CAAGCCAGIC ATGGAATCIT GCTGCAGGAC CICCCICIAC TACTICCTGI CCIAAAAATA GGG

SEO ID NO:1622: (Length of Sequence = 356 Nucleotides)

TTAATTITAA ASCAGATAAT ATTICAAATA TITICTITGA AATAGACCAT TIGICCIGCC TIGAAGTAIG TIAGTACATT
TTAAGAAAGT CAGIGGGITA AGGAGICAGT GCIGITAGTA TICATGCTTA AAACACTICC CTICIACCTA CCCTAATAAA
TGAGGGGCTC AAGAGAAATA TITICTAATIC TCTAGCGACA TGGCTAATTT TITITITITAA TGTATTITIG TATTITITAGT
ACAGATGGAG TITICACCATG TIGGTCAGGC TGGTCTCAAA CTCCTGAGCT CAAGTGATCT GCCTACCTCA GGCTCCTGAG
TCACTGAGAC TGTAGTTGTG TGCCACCATG CCAGGT

SEO ID NO:1623: (Length of Sequence = 361 Nucleotides)

TITGAGACAG AGTOTOGOTO TITTOGOCCAG GOTOGACTIC AGTOTOATTA TOTOAGOTOA CTGCAAGOTO CACCTOGOGG GITCACGCCA TITCICCTGCC TCAGCCTCCC GAGTAGCTGG GACTACAGGC GCCCGCCACC ACGCCTGGNT AATTITTTGT AITTITAGTA GAGACGGGGT TINACCATGT TAGCCAGGAT GGTCTCGATC TCCTGACCTC GTTGATCCGC CTGCCTCCGN CTCCCAAAGN GTTGGGATTA CAGGNGTGAG CANCCGTGCC CAGCCGINAA GTTAAGATAT TTTAAAAANA TCTCTGCAAG TTGAGGAAGT NTTTCAGGAC TCTTTCCTGC TTAGTCTCAC T

SEO ID NO:1624: (Length of Sequence = 350 Nucleotides)

CITTGIGAGO TITTIGACCI GOGGATCOG AGCCAGATIG ACAACAATGA GOCCIACATG AAGATCOCIT GCAATGACIC
TAAAATCACC AGIGCIGITI GGGGACCCCI GGGGGAGIGC ATCATOGCIG GCCATGAGAG TGGAGAGCIC AACCAGTATA
GTGCCAAGIC TGGAGAGGIG TTGGIGAATG TTAAGGAGCA CTCCCGGCAG ATCAACGACA TCCAGTTATC CAGGGACATG
ACCATGITIN TGACCGCGIC CAAGGACAAC ACAGCCAAGC TTTTTGACTC CACAACTCIT GAACATCAGA AGACTTTCCG
GACAGAACGI CCTGTCAACT CAGCTGCCCT

SEO ID NO:1625: (Length of Sequence = 333 Nucleotides)

GRITICIGIG AGACAAAGAA ATTATAAAGA TGGCAGAAAT TATTAGCGAC GITCIACCIC TATAATTCAC GITCCATGAA
TCAGTACITC ATTICITTIT TATGGATGAA TTAATATTCC ACTGIACAAA TATACCACAT CITGITTITC CATTCGICIA
GGITAAAAAAA TTITTATTIT TATTITTATIT TITTIGIAGA GACGGGATCT CACTGIGITG CCCAGGCTGG TCTTGACCIC
CTGGGCTCAA GIGATCCTCC CACCGTGGCA GTCCAAAGIG GGIAAACTGT ACGCTGGTCT GAAAGACCTT GCTGAAGAGA
GAAGAGGCAA GCT

SEO ID NO:1626: (Length of Sequence = 314 Nucleotides)

GACTGICCGI GGACACTGGI TITTAAGCCC AAGAACTGAA TATACAGTAG CAGTGCAGAC TGCCTCAAAA CAAGTTGATG GTGATTATGI TGTGTCTGAA TGGAGTGAAA TTATAGAATT CTGCACCGCA GACTATTCAA AAGTTCATCI AACACAATTG TTGGAGAAGG CTGAAGTGAT TNCAGGACGC ATGCTTAAGI TITTCTGTTTT TTATCGTAAT CAGCACAAAG NATATTTTGA CTATGTTCGG TAAGNTTCAA AAATATATAG TGATTTGTTT TACTAAATAT AGTTTCAAAT TCTAGGCTCA GGGT

SEO ID NO:1627: (Length of Sequence = 375 Nucleotides)

CCCTGGGCAC CIGGIACCIG GGGACCIACA AGGIGGIGAG GGAAGGGIAC GAGIACATIC CIINTCCCIC TGACCIGGGC
GCTAGAAGGG CAAAGAACCC GAGCCIGCCA GCIIGGCCIC CICCCACAGC CICCCTCGGA GGCAIGCCAI GCCAAGCACT
CTITCIGICI CIGITCAIGA ATAAAAGAGA TGGAIGGGCI TATICITATA GAGAAGIGAA TITCACITAC TCCCCTGGCC
CGAAAACIAG ACCAAATGAG GAACIGITIT AGCICATCAA ACIGITATAT TIATITICAA CAATGAAAAC AACACAACAA
AGIGGAGICA ATCCACIAAT TITITITAAAT CIAACACAAT TGITIGCACA ACAAT

SEO ID NO:1628: (Length of Sequence = 434 Nucleotides)

TECACAGECA CACCICCACT CITTATATCA TITTCICCAT CITTCATTIC CCATCIGIAC CICCAAAATT TIGCIATEAA
TCTAATTCAT CITTGCICIC TCTCICTCAT GGGIGCCTT GCTTCIGCCA GICTTCITC TCCIGCCCCA CCCAAACTIC
ATGAATTAGI CITTCICCC AGGAGCTCIG ATTTCTAGAC TGCTTTGAAA ATGCTGTATT CATTTTGCTA ACTTAGTATT
TGGGIACCCT GCTCITTGGC TGTTCTTTT CIGGAGCCCT TCTCAGTCAA GTCTGCCGGA TGTCTTTCTT TACCTACCCC
TCAGTTTTCC TIAAAACGAG NACACAACTC TGGAGAGTGT TAAGAATAAT GTTACTTGGT AATGTGTATT TATTGAGGAT
TGTTGTGCCTA AGAATCAGTA GGTTAAAATA GGGG

SEO ID NO:1629: (Length of Sequence = 341 Nucleotides)

CTCAAAGCT GCAGGGAGET GGGGGTGGCC CGCAAAGTGT TTGCCCGGGCC CCTGACTGTN TCCTTCCGGA GCTGCCGAGG ACTGCAGGAGA GGGCTTGTT TGGGGGTGGCC CGCAAAGTGT TTGCCCGGCC CCTGACTGTN TCCTTCCGGA GCTGCCGAGG ACTGCAGAGA GGGCTGGCT TGTCCCCTCT AGGAGCAGCT GGGNNGGTGT CTTGCCTGCA TCCCCCTTCA ATGGTTGAAA ATAATGATTC CACTTGTCAT GAACACCATG AAGGTATCIT GGCAGCCAGA GTCACTCCTG TTCCCGAAGT GGGAAACCTN GGGAGGGTCC TCAAAACCCC T

SEO ID NO:1630: (Length of Sequence = 380 Nucleotides)

CATAAAACCA TCCTACGATG TGCTGCTGCT GCTGCTGCTG CTAGTGCTCC TGCTGCAGGC CGGCCTCAAC ACGGGCACCG CCATCCAGTG CGTGCGCTC AAGGTCAGTG CAAGGTGCAC GGGTGCATCC TGGGACACCC AGAACGGCCC GCAGGAGCGC CTGGCTGGGG AGGTGGCCAG GAGCCCCTG AAGGAGTTCG ACAAGGAGAA AGCCTGGAGA GCCGTCGTGG TGCAAATGGC CCAGTGACCC CCAGACGCGG AAACCGGGTG GCAGGGCCCAG CCTGGGCCCA GGCATGGAAA CGGACAACCC CTAATCGCCT TAGCTACTGC TTCTAACAAC TCTTTTCCCT TGTGTTAAGG GAAACCAGGT TCAAGGGGGG

SEO ID NO:1631: (Length of Sequence = 383 Nucleotides)

AGAGGATTIA TITIGGACAGG GCTGTGCTGA GAGTCCCACC CTCACCCCAC AATGGGCGG GGCACTGGCA TCGAACACCA
AGCTGAGTGA GAAGGCCTCC TCCAGGCCTC GCAGGGAGCT TGCTGGCTTC TCCTGGCTCA CAGCAGACTG GGCCCGACTC
CCATCGGAGG AAGGCCAGCA TCCTAGGGCA GCCAGTGGAG GGCTGGCAGA GGGCTGTGCC TNGAAGGTCA CTGTGCTATC
TTCCAACCAC ACTGTGTGAG TCTCAGATAC CATATGTGGA ATCTGCATCA GGAAGGTCAA CTTGAGGTCA TTTTAAAAGG
GATTCTTCCG GNAAAAGGAG CNCCGCATCG GGCGNCTTAA NCCGGCGTTT CGGTTCATCC CGA

SEO ID NO:1632: (Length of Sequence = 424 Nucleotides)

GGGAAGTGAG CTCCTGAACC AACTCTGAAG GAGACACCCA CTTGCTAAGC CAGTCTCACT CTAGGACACC TGCCTAGCGA
CCAGCAAACC TGGAATGAAA GGGCAAGTTC CTCAGTGCCC CCTCTGCATC AAAGGGAGTG GCTCTGCCCT CTCTAGTCTC
TGACTACCTG CTTAGTGATT TTTGCTTCTG TGCTCCCAGA CCCAAGAAAA CCACGTCTCT TTTCTTCCTT CATCGACTCA
TCCCCTTCTT ACCCTATATT GTCTCCTCCA CTTCCTGCCT CTGCTGGCCA GGCTTAAATC TGGGCCACCA GCCTTCCTGG
GACATACCTA TTTCCGCAAC TGAACCTTCC CAACCCCTAG GAAAACAAAG GTATTTTACA AGGCCTCTGG ACCTTGACCC
AAAGAGGCCAT GNACCATAAT TACT

SEO ID NO:1633: (Length of Sequence = 417 Nucleotides)

TTITITCIAC AGCATCITIT TATIGICITI ACCATTACIT TAATGCATIT TAAAATTIAT CIACATTAAT TGGGAACTAT
TTGCATTITI TICATCCICI CICICITIIN CITTINCITI TITIGGATIT GICTIGGCCA GAGAGGITCI CCAACACCCG
GGIGGACTIG GAATTITTIA TCAGCIGCAA TCIGAAGACT TGICTITACI GIGGAATAGG TGACATTCCI TIAGGACCTC
AGAAGCTCAA GIAGITTAAT GCCAAGICIT TCCAGAGCCI CACICICITI TATITITIAA ATTAGAATIG TGATTTATIG
AACNCITACC AIGGGGITCA TATAATTINI NAATNGANCA GCITTATIGA GGIATAATTC AATACCCCIT TAAAGNATGI
AACCCGIGGG TTTAGAC

SEO ID NO:1634: (Length of Sequence = 423 Nucleotides)

AATATCCCAA ATGIGCAATG CATCACCIGA GACAGAAGGC AGAAAGCATC AAGCICICIG TITATCCCAA TICAATGACA ACCAGAACIT ATTITITITIG AGATGGGGIC TCGITCTGIC GCCCAGGCIG GAGTGCAGTG GGGCATTCAT GGCTCATCGC AGCCTCCAAC TCTCAGICTC AAGCAACCCT CCTACGICAG TGTCCTGAGT AGCTGGAACT ACAGGCATGC ACCACCACAC TTGGCTCAIT TTTAAAAAAT TTCTTGTAGA GACAGGATCT TGCTACATTG CCCAGGCTTG AGGTGCCGTG GTGCATTCAC AGCTCACCGC AGCTCAAACT CTTTGGTCTC AAGCGATCCT CCTGNCTCAG CCTTCTGGGT GGCATGCAC CAGCCATACA CCACCATGTC TTGGTCAATT TCT

SEO ID NO:1635: (Length of Sequence = 384 Nucleotides)

CAAAACTCAC TITGACCCCA TTAAGAGGCA AGCCTGGCAC ATCTATCCCT GGGCCTTTAG AAAGCCATTT GCCTCAAATG
GCTATAGGGT TGTGGGGTGG AGGGAGGAAG GGCTGGGAGG GAGTNGGGAG GAATTGCTAG CTGTAGTGTG ACACATTGTA
GTGTTTGCCA GGAAATGAGC CAGACATGGT GGTGTATGCC TGTAGTCCCA GCCACCCAGA AGGCTGAGGC AGGAGGATCG
CTTGAGACCA AGAGTTTGAG CCTGCGGTNA GCTGTTAATG ACCACGGCAC TCAAGCCTGG GCAATGTAGC AAGATCCTGT
TNTCTACAAG AAATTTTTTA AAAATTGAGC CAAGTTTGGG TGGTGCATGC CTGTAGTTCC ACTA

SEO ID NO:1636: (Length of Sequence = 362 Nucleotides)

CAAAATGACT GACTACAGCA ATGCCTTCCG TGTGCCCCAC ACATCATGAG CACCGCAAGA GACAAAAGAT TAACTATGAA ATATAGTAAT CIAAGCAAGC CCACCATAC ATATTTTTGG GGATTTCCCA CCATCCTGAA TAGTATCACT GCAGTTGACA CAACTTCCAG GGAACTGCAG AGTAAGTGCT TAATATTATC CACGAGAAAG CAAAACTAAA TATTAGTGTG CACATTTCTG AATGAGAAAC TAATTGCTTC ATTGATTTCA ACAATGTAGT GGNAGNAAAC TATTTCAGAT CTCTACAATG CCTAAATGCA TTCTATTTAA ACTCAAGGTA CTATTTTCAT TTTTACCATA CT

SEO ID NO:1637: (Length of Sequence = 205 Nucleotides)

GGCGCCCGAC GAGGCTCAGA CCTCTINIAC GNCGACTACT ACGAGGACGG CGAGGTGGAG GAGGAGGCCG ACAGCTGCTT
CGGGGACCGAG GAGGATNACT CTGGCACGGA GGAGTCCTNA CACCACCAGA ATAAACTTGC CGAGTTTANC TCACTAGGGC
CGGACCCGTG GCTCCTTAGA CGACAGACTA CCTCACGGAG GTTTT

SEO ID NO:1638: (Length of Sequence = 253 Nucleotides)

CACTCAGGCT CACCGTCCTG CTCTCTGCAC CAGCCTTTCC AGAGCATNCC AGINCTCATG GCTTCATCTG TTAACTGTTG ATCACTTCAG TCCTGATTTT TAGACCTAAA TGGTTTCCTT AACGCCATTC TAACTGCCTG TGACTCATTT TCACTTACAG TGTTTATTGT AACGCCAAAC CAACAAATCA CAGGTGCTTG CTTCTGTCCA TAAATCTCCC CAGTCTAACT TTTTGTCATT CAACATGRCT CGT

SEO ID NO:1639: (Length of Sequence = 360 Nucleotides)

TGTGGCCAAG GACCCTATCG TCAATGTATG GTACTCTGTG AATGGTGAGA GGCTGGGCAC CTACATGGGC CATACCGGAG CTGTGTGGTG TGTGGACGCT GACCTGGGACA CCAAGCATGT CCTCACTGGC TCAGCTGACA ACAGCTGTNG TCTCTGGGAC TGTGAAACAG GAAAGCAGCT GGCCCTTCTC AAGACCAATT CGGCTGTCCG GACCTGCGGT TTTNACTTTG GGGGCAACAT CATCATGTTC TCCACGGACA AGCAGATGGG CTACCAGTGC TTTTTTTGAC CTGCGGGATC CGAGCCAGAT TGACAACAAA TGAGCCCCTA CATGAAGATC CCTTGCAATG

SEO ID NO:1640: (Length of Sequence = 321 Nucleotides)

GIGGGACGCC CTCTGCCTTG TCCTGAGAGC AATGTCTTCT CCATGGGGCA GCATNGGCCC TGGATGGGCC TGAGCATAGC
AGACCACGTG GTCACATGTG CATGTGTGGA CATGTGTGCA TGTGTGGATA TGTATGCTCC TGAGTGTATC TGCATGTCCT
NCCTGCACAC ACAGTGCTCC CCTCCGATGC TGCCAGCCTG TGGTGGACTT CCTCTTCTGA CCCCTTTCTT GCCNCCGGNC
TGTTTTATCA GTGAAAGGAC TTAACTAAGC AGATCTCCAG GTTCACCTTN TGGAACTCAG CTCAAGGTNA GCACAGCAGG
T

SEO ID NO:1641: (Length of Sequence = 256 Micleotides)

GGIGGIGCCA CIGICGIGAT AGITTITCCC ATCITAGIAG CCGNACCCAT AAITAATGCC TACTCACATC AAGITAGCAC CACTCAAAIG TGGGCCATTC ACAGGCAGCC AGGGATCCTC TTGGNCCGTG AGGTTGGGGG CTTNCATCAG AATGCAAATC

TROCGAGGOG TGAAGCACAA TITAKTICAA CIGCOATKIK TIOCITCACA GIAAGROCIT CIGGRGGAAG GAAGCAGIGI GIITGAGITA TACCITAGGO CAAGCI

SEO ID NO:1642: (Length of Sequence = 295 Nucleotides)

AAAAGCCCCA GCCTCAGGAC CCCGGTCACA GGCACCCGGG GGTGGGGGTG ACCAGCAGCA GTTCAGAGGC AGGTGTGGGC
AATGTGGGCC TGAGTCTCCT NCCCACTCAC GTCACTNCCC GCGGGGACAC AGCGGCATTT NTGGGGCACT NGGCATGCCG
GGTTCCTAAC CTCAATTATT CATTCTGCTC TCAGGCACCT CCTGACGAGA CCCTGGCCCA GGAGAGCTCG GCTCGGGGAC
AGAGGAATGA GACTCAGTGG GACGCAGAGN CCAACCCCAT CCCCACCCCT GGGCT

SEO ID NO:1643: (Length of Sequence = 359 Nucleotides)

ATCATIGGIA GITTAAACIT TICATCIAAT ATTAGATIGC ATGCAGGATT TIATATCIAA TIACICIGGC AGATGGCCIT
TAGAAAGITC AAAAATAAAA TGCAGCAATT CATATTGGCA GATTTACTAT TGAGACCAAT GCTTTCITAA CTAAAAGGIT
TIGITTAAAA TCGITAGITT AGGAAATCIG ATAAAGATTT TIGAATATCA GAGCGITTAA AAGAGATTCT TACITTACAT
CTGGCATATT TCTTGIGITA CATATTATAA TTCCATTGGA ACATGGCIGT CIGIAAAACT ATGTATATGA TCCGGAAGAG
ACTCAAATTA AATTAAGGIT TAACAGCCAT CAAGITCAT

SEO ID NO:1644: (Length of Sequence = 293 Nucleotides)

TGAACCCGGG NGGCGGASTT GCAGTCAGCC GAGATGGCAC CACTGCACTC CAGCCTGGGT GACAGAGCCA GACTCTGTCT
CAAGAAAAAA AAAAGAATTA AAAGATGTGA ACAAAAGCAA GAAAGTGCTG TATGAACGAA ACGGAAATAT CAATGAAGAG
AAATAAAAAT TATAAAATTC AGGAAATGAG ANGTACANTA NCAGNAAATT CACTGGAGAG ATTCAAAAAGC ATATCTGAGC
ACGTAAAAAAA AGTAGTGAAC ATGAGATAGG TCAAGGGAAA AGTACTGAGT CTG

SEO ID NO:1645: (Length of Sequence = 332 Nucleotides)

AAAAGCIGGA TATTAGGAAA TGTGAATATT AATTCTGAAT TTGTTACTGA CTCAGGATGA CCTTGCATGA TGCATCCAAC
CTTCTTTTCT CTATATCAGA AAACTAAAGA ATAAATGTAA CATCACATTC TTTTCTCCTT TGGGACAAAC AACTATGTAC
AATTGAATAA AAATGAAATT GCATAAGTNG TGGATAGAAT ATGTTTGGGT TGGTTTGAAC TTAGCACACT GTTTAATAAT
TCAACATTTT TTATACCTGT GCAATAAATT TTTAAATGAT GTCTGAAATG CTTTGAAATC TTCAGAAACA GGTTTATAAA
TCGCATAAAA AA

SEO ID NO:1646: (Length of Sequence = 210 Nucleotides)

GAAAGINCTC CCAATCACTC TCTGCACAAT GAAGTGGCGG ATGACTCCCA GCTTGAAAAG GCAAATCTCA TAGAGCTGGA AGATGACAGT CACAGCGGAA AGCGGTGGAA TCCCACATAG CCTGAGTGGC CTGCAAGATC CAATTATAGC TCGGATGTCC ATTTGTTCAG AAGACAAGAA AAGCCCTTCC GAATGCAGCT TTGTTAGCCA

SEO ID NO:1647: (Length of Sequence = 246 Nucleotides)

TCCACTCCAA GGGTTTCTGA CCCAAGAGGT GGGGACCAAA ACCATGCATT CCTAAGAAGT CCCCAGGTCA TGCTGCTGTT
GCTGGACTGA GGACCACACT TTGAGAACCT GTGCTCTAAG TGAATACTTG GAAGTCGTTT CAGGACATGG GGCATAGAAA
CTNAGGAGTA GCTGAGAGGA AAATNAAGAG AAGCTGAGAA GAAGCTGAGA ATCCTCACAG GAGCAGACAG AGAAATGTGA
AGGGTT

SEO ID NO:1648: (Length of Sequence = 338 Nucleotides)

TCCACTCCAA GGGTTTCTGA CCCAAGAGGT GGGGACCAAA ACCATGCATT CCTAAGAAGT CCCCAGGTCA TGCTGCTGTT GCTGCACTGA GGACCACACT TTGAGAACCT GTGCTCTAAG TGAATACTTG GAAGTCGTTT CAGGACCATGG GGCCATAGAAA CTGAAGGAGTA GCTGAGAGGA AAATGAAGAG AAGCTGAGAA GAAGCTGAGG ATCCTCACAG GAGCAGACAG AGAAATGTGA AGGGTGGGGT TTTATGTNTG GGAAAGGGAC CCGAAGCCCA GGCTGAAGAG TTTTAACTTT GGGCCCAGAA ACTCAACCAT CAATGGAAAC AGGGCAGT

SEO ID NO:1649: (Length of Sequence = 275 Nucleotides)

GCACCTINAG GATIGAGACC CGGAAGGCTT CAAAGGCTGT CGCAAGGAGG AAGAACTGGA AGAAGTTCGG GAACTCAGAG TTTGACCCCC CCGGACCCAA TGIGGCCACC ACCACTGTCA GTGACGATGT CTCTATGACG TTCATCACCA GCAAAGAGGA CCTGAACTGC CAGGAGGAGG AGGACCCTAT GAACAAACTC AAGGGCCAGA AGATCGTGTC CTGCCGCATC TNCAAGGGCG ACCACTTGGA CCACCCGNIG CCCCTACAAG GATAC

SEO ID NO:1650: (Length of Sequence = 270 Mucleotides)

AAAAGCCAGA GGGATGAGAA TGAGAAAGIT AAAAGGGAGG TCAGGAAAGC CATCITITAG GAGAAATATA AATMGACAAT
SCITTAAAAA AGGAGCTGCC ATCATATIAT ACCCTGACCC AGCTGGATAC GAACAAATTC AGCCTTGGCA ATGCAAGTCT
TACATCIATT TIATATAGAT TGIATAAAAG AGAACTGGAA GCATTTTCAA GAGGGGTATG TATGTGTTTG TGTGTGTCTG
GTAATTAATG AAAGAGAGCC TATTGAATTT

SEO ID NO:1651: (Length of Sequence = 372 Nucleotides)

TCTTGCTTTT TAATTGTATT TCTTAACACT AGAATTTCT ATTTCAAGIT TTTGTACGIG GCCTTGCGTC TCCTTAGTAC
ATTTTATAGT CGCTGTAAGT TGATTCCATT TTTCTTGAAA TTGAATTCTC ATCTGACCTA ATTTCTTCCT TGAATCCTAC
ATCTCACTTT CTCAATGGAC GCAGTGACGC AATGAAGCAT CCAGCAAAGC TTTTGTTGTT GATTGTTTAG GACGTCACCC
TGTTTTTGTT GAAGTTGTCT CACAACTACT TCTCTTTCTG CTTTCTCTCT TTCATATTGA CATTGTTTTT CTTTTCAAAT
GGATTAACTT TATTGATCAT CCTCTTGTMC TTCTAGCAAA AGACGGGTGC TT

SEO ID NO:1652: (Length of Sequence = 314 Nucleotides)

TTTCTGAGTA TGCTGCACTG GATTATTAGC ATGITAAATA GTCAAAGGGA CTGGAATAAA CATCAGGAAG ATTTCATAAA GTGGTGTAAG TAGAAAAAAA AGGTTAAACA ATGAGCTGCA TGTTGATAAG TATAAGACAC TGATCCAAACC CTCATTTTAA GAACCATGAT ATTACTTAAN CTAGAGTGTT AAGGTCAGCT TAAGTCAAAA TAAAACAAAG CTTCCAAACC CTCATTTTAA ACCACAGTAGA TAATAGATGA NTCTTGTATC TTGGGAGATA GTACAAGCCA AANGTTACAG CTGTGTTAAA ACCT

SEO ID NO:1653: (Length of Sequence = 323 Mucleotides)

TAGATATGAT GGCTGGAGCT GCAATAGCTA ACTTGCAACT ATGAGGAACT ATAGGACTTT GGTCTTAACA TTCCTGAGCT
CCTGAATCAA TACTTTAACT ACCTTCTATG AGACTTCTTG TCACATGAGA AAAATTAAGC CCCAAATTAA ACCCCTGCCT
TINACTGTAA CTCTCAATTG AGCATAATTC CTAAATGNTT TAATCAATTC TACTCTACTC TGGCATGATT TINAAGGCAT
TAACCATAAT TTCCTTCCAA TCTAAAAAAGG GAACTANTAC TTACTGGAGT ATCTAGTATA CATCAGATAC TGTGTATATA
GGC

SEO ID NO:1654: (Length or Sequence = 352 Nucleotides)

ATCTTGGCCT GCAGGAACAT GGCAAGGCG AGTGAAGCAG TGTCACGCAT TITAGAAGAA TGGCATAAG COTAGGTGGA AGCAATGACC CTGGACCTCG CTCTGCTCCG TAGCGTGCAG CATTTTGCTG AAGCATTCAA GGCCAAGAAT GTGCCTCTTC ATGTGCTTGT GTGCAACGCA GCAACTTTTG CTCTACCCTG GAGTCTCACC AAAGATGGCC TGGAGACCAC CTTTCAAGTG AATCATCIGG GGCACITCIA CCITGICCAG CICCCICCAG GGATGITTIG GIGCCGCICA GCICCIGCCC GIGICATIGI GGGTCICCIC AGAGICCCCA TOGATITACA GG

SEO ID NO:1655: (Length of Sequence = 325 Nucleotides)

AGGGTAAATT GTGAGACTGT TTGTATATAT TNTTTGTTTA TATGTTTTTG TTGTTGTTAT GTTGTTATNT TTATTTATAA
AATGATAGAT CTGTGGGTAG GTTCTGAGAA ATGAATAGCT TGTATTTCCT TTTTTATGAA AGAAGAACAA AATGAAGTTC
AAGTGGAAAG TATCTCCAGA AAGTTTAACA TTTTCTTATT AACCAACTCA TTGATTGGCA TGTGAAACTT GAGATATTTT
ATATAGCACT TTTTAAATGA GGATCTAGCT TCACTNTATC ATACAACCAC ATTTAAAATA GCCAGGTCCA TGGTCATTAT
AGGGG

SEO ID NO:1656: (Length of Sequence = 285 Nucleotides)

GAGENTIAAT AGAATAGATC AAAGCAGAAT GCAGTGTGT CATGTCATAG GTTGACTTCT CCAGGAAAACC GACCCCAAGT
GGAAGGTTTA CATGCAGGTG GTTTATTAGA GAGTGATGTT GGGAAGAACA CCTGTAAGGN AAGAAGGGAG CCTGGGAAGA
GCACNOCNAG AAGGTGAACT CTGATTCACT TGCAACAGAG TCCTAGGCTG AGTGCATGGG ATNCTGTAGA GTTGGGGATG
GACCTTCAGA GATATTCCAA ATAGAGAAAG AATTCCTGTT TACTC

SEO ID NO:1657: (Length of Sequence = 385 Nucleotides)

GACTIGACIT TECTITITIC CCCCCAAGIA GAACTAATEC TAGCITCCAG CTIGAAAGIA AAACTCCAGI GIGGAGIGAA
TITIGIGICT AATTATAAAC CIGTAACCAA AACTCAGACA TCIGGIACIG GICTITIGCAT TGAGATIGGI CCCTGIAAAA
CCCCCTTTAA AAGCATATIG CATTIAGIAC AGAGCTCTIT TTIGAAATGN AGGCTGGAGA TGIGCATTIT TCACGGIGIT
AACTGGITGI ATCITATIAG CAAGGAGATT GGGGGTTTITG AGTGTTIGCG TGGGTGGGTT TCAAATTIGC CAGGGGAACC
AGTGGGCAGG CTGCTAGCAA GGCAGTGAGG AAGCTCTIGG CAGCCAAATG GGGTGCATTI CAGGG

SEO ID NO:1658: (Length of Sequence = 338 Nucleotides)

GATCAGGACC TCTTCTTCCT CCCAACACTG CCCCAAGAGC CCGTTGTTAA ACGTTTACCA GCACACTACT GGGCTGTTC
TCTACCACTT GATTGAAATG ATCCTTATGG AAGCACAAAT GACTTCACTG TCACTAAATC CAAGGGACAA TTTTTAGTCT
CTATTTTTCT TCAACTCTCC AGGATGTTTG AGAGCTGATC TTTCCCTCCC TCTTGAGCCT CCTCTCCTGC CTGGCTTTTA
GGGGTCTCTG CTGACTTTTC TTCATTTCTA AACACATGTN CTCAGGGGGT CCTCAGCCCT GCAAGGCCNA TGCACTGGGT
ACCCAGTCCT GTGGGCCT

SEO ID NO:1659: (Length of Sequence = 346 Nucleotides)

AGIATGIGAA GICAATCACT TITITATATGC AGATAATATG CGACTIATAA TGGAAGGICA CGITICAATA GCAAACAAAA
AAGCIATAAG TAACAAAGAA TAACAAAACT ATAAATGIAT AGGCTCIACA TAAAGAAAAC TATAATICCA TAAAGGATCT
AAAATAAAAC GAGIAAATGG AAAGACAAGA TGTGTTGTGA GATACGAAGA ATCCATGATT AAGITAGAGG ATTCTTGGAT
GACAGTAGAG TAGAAAGCAC CAAGAATGAG TCTGTATACC CAGAGAACAC TTACGCTGGT AGGAATCTAT CTCATACAAC
TATTATGGAG CTCTCAAAGT ATACTG

SEO ID NO:1660: (Length of Sequence = 240 Nucleotides)

GATAGAATAG CCAGCCTTCC ACTTGAATGC ACTGCCATAT TGTCAAGCTG CATTCCTTAA GCATCACTTC TTAGAGGCCT CAAGCTTCTC GCGAATGTTT GATGACTTAA AGGGGAAATG AACAGGTTGC AATNATGCTT GTCAAGNTTC TEXTTGTGAA CCTCTATTTG GACAATTCAC ACAAAAAAAG AAAGCAGGTC ATTTTCTAAT TCAGGATATT ATTTCTTTTT AAAACTGGTA

SEO ID NO:1661: (Length of Sequence = 294 Nucleotides)

AGCACCTCCC CTGAGGGCCA GGCCTTGGAG AACCGGATGA AGCAGCTCTC CCTACAGTGC TCAAAGGGAA GAGATGGAAT
TATTGCTGAC ATAAAAATGG TGCAGATTGG CTGATTCATC CTGGGCCCTG GCCGATATGC ATATCAACAT TTATACATGG
AACTGTGAGA ACATTKTGCC AATAATCATT TAATATATGC CAAATCTTAC ACGKCTACTC TAAACTGCTC TAATGAAGTT
TCAGTGACCT TGAGGGCTAA AGATINITCT TCTGGTGTAA GAGCTCTTTG GGCT

SEO ID NO:1662: (Length of Sequence = 291 Nucleotides)

GATTITCATE AGGCAAATNA AAGTAACCAC AGAAACAATT CAGTAATACT ACTAAGAGAG ATTAACTTCC CACTGGCCTT
GGAATAGCTA AGTGCATTGA TTTTKGTGTA GITGTGAGTT TTTTTCTYTC ATTGATATTT TACGTATTKC TGGGGTAAAT
GTATTTTWA CATGCATTGA ATGTGTAATG ATCAAGTCAG GGTATTTGGG GCCTCCATCA CCTTGAGTGT TTATCATTTC
TATGTGTGGT AACATTCCAA GCCCTCTCTT CTAGCTTTGG AATATATAGT G

SEO ID NO:1663: (Length of Sequence = 345 Nucleotides)

GGCAGTGGGA CTCTCTGTGG ATAGACTGAT TCTTGTTTAG AAACAACAGC AAAAAGAAGA AGGCAGGAAA GAAACTCCCC GGCTCGGAGG AATGTCTCTG TGATCCCCAT TCTTGATGGA GGGAGTGAAA AGGGGCCTGG NCTTGGCCCG CTGCTCTCCT GACAGAAACA GTAAGTNACA CCAGGACAGA AGGCAGGAGC CCTGAGAACT CACGGCGCTC TGCATGGTCT CCAGCCNNNC ACCCGTCTCC AGCCACCCCT GGAGCGGCCG TGGGGAGGCG GCAGAGGGGG CTTTTCGGAG GGCCCACTAT TNCCACACGT CTTTCTTING ACACCCAGAA AACTT

SEO ID NO:1664: (Length of Sequence = 334 Nucleotides)

GIAAATAAGA AAGIGAAATA ATTCCIATAA TGIAAGGIIG ATAGAAGATA ATCATCAGGG TCAGAATIAA GAGGICIIGI GGITTAGGAA GCATAAAATT ATGIAACTIA TIGITTATIT CACICAGAAA ATAAAAGIAT TAATGAAAAG AGITAGAGAT GAACAGATIG ATACAAACTG TICIATGGIT TACAGCTIAA AAAATAAAGG TACATTIAAT GCIATGCATT TIGAGAATAA TGICITTTAT GCINITCCIT TITACATATG TATCINITIG TATTIAAGGI CAAAATAGAT TGACATTACT AATTACITCA CTATTAATAA TIAA

SEO ID NO:1665: (Length of Sequence = 310 Nucleotides)

TGTACINCTA TGAAGCATCC CTTCCACATC AGATCAAAGA CATCTTAAAG CCAGAAATAA TGGAGGAGAT TGTGATGGAA ACACGCCAGA GGCTTTTGGA ACAGGAGGGA TAAGGAGGTG CTCCAGAAGC ACGGGACTNT GGACCTTGCA GGAGTGAAGA CTGTRATGTG TGGTCCCCAT ATGTGGCTCA GCAAAGACTC GAGAGATCAT CCCTTTGTCT GCATTGACGG CCCTGTGACG GCCTCCAGCC CACAGGCCTG CTTTCTCCTG TCCTAACACC AAGCCTGGGT GGCAGATGAA CAGTGCTTCC

SEO ID NO:1666: (Length of Sequence = 352 Nucleotides)

TTITITITIA CATACAAAGI TIGGATITIT ATIGAAATCI TGITAGGIAT CAAACAAATI CIGCITTCIT CAGATAAAAA TATICICICA GATGICICCA GATAACIGCI AAGICIAAAT TGGICCITCA ATGICITATI TITATIGICC TCGIGAAATG TTCATATACA GITAAGATGI TCCCCAAAAGG ATITITATCG TGIAAAGGAG CGIACATGAC GACCICTACC ACTGCCTCCA CTAACAAACT TTCCTCTTGA GCCTCCACTG CCGCTATTTG CACTAGCCCA GGGAAGGTCC AAGICCCCCA CGACCTCTAG AAGCACGGTT CCGAGGGACT TTGGCGGTAA CC

SED ID NO:1657: (Length of Sequence = 287 Micleotides)

GACAATNATG COGCTGCCCA CATTITGGTC CATTCTTTT TITATTATGC TICTCTTNCT TOGACTGGAT AGCCAGGGAT
GTTTCANCTT CTCGCTCGTC AAGTACGTAC CCCTGACCTA CAACAAAACA TACGTNTACC CCAACTGGGC CATTGGGCTG

GECTIGGAGCC TIGGCCCTTIN CTCCATGCTC TIMENTCCCT TIGGTCATCGT CATCCGGCCT CTGCCAGACT GAGGGGGCCG
TTCCTTTGTG AGAGTCAAGT ACCTGCTGAC CCCAAGGGAA CCCAACC

SEQ ID NO:1668: (Length of Sequence = 300 Nucleotides)

CCAGACAAAT ACCAAGTITA TITCACAAAC ACTAGGAAGA TGGGTTGAGG GTGGAGGTGG GGGACACAGG TGCGCANTGC ACAGAGTCAG CAGCAGCAGC CTGNTCCCCG CACTGAGGAC TCGGCCTGGA CTGCAGTGCC TCCAAATCAA CACGCAGCAA GAGGGGAGTIN CAGNGAGGGC CCTNAACACC AAGCCTCTGA AAGGCTAAGG GACACAGCTC CATCTGTCCC AGGAAAACCA GCAATAAATA AAAGTNNGGC ACGGCCCCAC CCACACATAT CATCTAGTCA CCCATCTTCA

SEO ID NO:1669: (Length of Sequence = 334 Nucleotides)

TTITAATGAC AGATTITCCI AAAAGAAACC ACTATAACAT CIGICCAAGI ACTCCAGAGA AAACAAAAAA TACATAAAGA
TTAAAAGICT ATTACITTAA CAGCACATIG CCAAACACG ACAACTAGGA TAAATGCCAA GAAACCTTAA AAAATAACIT
TAAAAAGATGC AACGITCAAG CCATTCAAAC GCGTAGGITC CACAAACAAC AGGNNAACAA GICCAAGAGC AGITCIACIT
GTGCATGATG GTAACTCAGA CTGTACITCA TCAAAGITCA TTCAGGIGIT TCATAGGCGT CIGAGCAGAG TTITGITTIT
TTCTITCCTT GCTT

SEO ID NO:1670: (Length of Sequence = 287 Nucleotides)

GATAAAAGAG AAACAGCGAA GTTTCAAGAG AAAAACTTGA GGTCTTAATA ATTNITGGGC AACTTGACAG CAGAACAGGG
TAAAANTGAG TTAGCTACAA AGGCTCATCA GAAAATGGCA ATAGATTCCA GAGAGATTTA ATAACTACTT ACAAACTCTG
CTATAGGTGA CAAATCTGAC CATGATAAAA GCACCGTAAA TGATATAGGT AACACTGNGC ATATGAAAAC TCAGACTGTG
CACTAGATAA AAAGGAANCC CAGCATACAG TGTTACCACA TGTAAAT

SEO ID NO:1671: (Length of Sequence = 187 Nucleotides)

GATAAAAGAG AAACAGCGAA GTTTCAAGAG AAAAACTTGA GGTCTTAATA ATTTTKGGGC AACTTGACAG CAGAACAGGG TAAAAWTRAG TTAGCTACAA AGGCTCATCA GAAAATSGCA ATAGATTCCA GAGAGATTTA ATAACTACIT ACAAACTCTG CTATAGGGTG GACAAATCTG GCCCATG

SEO ID NO:1672: (Length of Sequence = 329 Mucleotides)

ACATCACAAC ATCGITTATT ATGIGAATIT TITACAATAC AAACAAAAAA TACAGAAATG CAATATATGA ATACAGCITAA ATGCAGAATG GIGACITTIT TCTCTTCAAG AGGCCATGAT TCCCATTTCT AGTAAAATAA AGAGACIGCA TATAGGTAGA AACAGGITGG TCATTAGCIT CACAAATTTTG CCTAGAAATG ATCTATAAAAT GCATTTCCCC CCCTGCTACT TACCCTAAAG TGTAAAAAAGG GAGITAAAAGG AAAGTTTCCT TGTTGGTTCC TACCATATGA AAGATGCTAT ATTCTATTTT AGCAGTGCCA ATATATGGG

SEQ ID NO:1673: (Length of Sequence = 386 Nucleotides)

CTCCCTACTG TGATTCTCAT CAAGCTGGAA GCGINGTGAG AAAGCACTTC AGTTTCTTCC CTCGGATATG AACCTGAGCT CTCTGATGAG GTGGTTTAGA AGTGGCCCTG GGAGAAGCCC ACTTCTTGGT CACAAGATAC TGCATTCTCC TGGCAGATGA ACCAGCTGCT TCCAGCATCC TCTGTGTGGG TCCTCACGCC TAGCTGCTCT ACGTGCTGGC TGCACAGTGG CATCACATGG GGAAGTAGAA AAACCTCTGA TGCCTGTCCC CACCCGGCTT AATCACAGTG AGGACAGTAG AATTTTT TTTAAAGAAT TGCAGGGGCC AGGGCGTGGC GGGCTTCAGA GCTTCTTAGC AATTTTT

SEO ID NO:1674: (Length of Sequence = 377 Nucleotides)

SEO ID NO:1675: (Length of Sequence = 381 Nucleotides)

CAGAAGTCAA TCAGCTACGC ACCCAGTTCT CAAAGACCTC ACATGCTAGG GAAGGTGCGG AGGCAGAGTT GTGGTTCAGA
AGCAGTTACA GGTCTCAAAG CAAGAACAGC AGCCAAAGCT TCCACGCCCT GACGCTGCCT CTGAATGGTA AACCAATGGC
ATATGGTATC CACAGCTAGG CTTTGCTTTT TTCTGAGTGA AGGTAAAAGG CATTTGAAAA TAAACCAAAG TTTCACAGAC
TATGTTTATG GAACAAACAT GGGCCATTTT CAGGGATATA AAAGTCGATG TTCTATGTAG GCCCCCATAT GAGTATTTAT
CTACTTTTTA TTTACTTTAT TTTATGGAAT TTATTTGNCA AGGGGCTTCA CTCTGTTCGG A

SEO ID NO:1675: (Length of Sequence = 404 Nucleotides)

CIGIGITICAT IGCITGAGCC CATCACAGIT TAGCICICAC AGCITTAATT TACTAGCCCA IGAGAAGICA GCITCAAAGA
ACACCATITC GACICICAAA GAACATIATC AAIGIACATG GATAGCITCC AACITCATAA GGIGITICIC ICTACCIAGA
GCAATTAACA ITAATITIGCA GAATAGIGIT TATIGAAAAC CITTGIGIAT CICCAACAAA GIAATAGIGI ATIGATITCA
ITCCIACIAT CITCAACIGI ATCATTAAGA GGAATITCIT AGGIAAGICI ATATGCAGIA AGCAAGIAAG ATCGCAGAAC
ATCAAAGGGI GGAAGIAAAT CCCAAAACIG GWITTTACCT TCCITTCCCT TAGGIGAGGG AAAGGAATIT ATGGITTTAA
AGCT

SEO ID NO:1677: (Length of Sequence = 388 Nucleotides)

ATGGACAACT ATGAGCCAGG AGTCTACACA GAGAAGGTTC TGGAAGCCAC TAAGCTGCTC TCCAACACAG TCATGCCACG
TTTTACTGAG CAAGTAGAAG CAGCCGTGGA AGCCCTCAGC TCGGACCCTG CCCAGCCCAT GGATGAGAAT GAGTTTATCG
ATGCTTCCCG CCTGGTATAT GATGGCATCC GGGACATCAG GAAAGCAGTG CTGATGATAA GGACCCCTGA GGAGTTNGAT
GACTCTGACT TTGAGACAGA AGATTTTGAT GTCAGAAGCA GGACGAGGGT CCAGACAGAA GACGATCAAC TGATAGCTGG
CCCAGAGTTG CCCCGGGCGA TCATGGCTCA AGCTTCCCCA GGGAGCAAAA AAGCCGGAAG ATTTTCGG

SEO ID NO:1678: (Length of Sequence = 428 Nucleotides)

TAACTGTGCA AATAATCCAT GAATATATTG TTTTTATACA GCATTACAGA TAAGGCTTGC AGCTCTATAG ATCACCCTCA TCCACTCCTT CACTCCATTG CTACACTTAA AAGCCTCACA TGCTCTCCTG TCCTCTCCAA AGGCAGCTGC TAGCATCAGC GCCCACAGTA GCCTTCTTTT GTTTCCTGTT TATAAACCAT ACATTTTCTA TGGCTACACA TACGTGTATT GTTTGATGCT TTCTAATAAA ATTGTATCAT AGTGGTACAC ATCTTTCACA CTTTCCTNAT TACAGTCAAC ATTTGGNGGA ATACAGAATG CAGCAGATCA AGGANCTTTT CTCAGTCTTT TCTAACATCN CCCCAAATAC AGCCTCACTA TGGGGTCCAT TTAGGNGGCT CATTGGTTTT CACTCTACA ACGGTGGC

SEO ID NO:1679: (Length of Sequence = 256 Nucleotides)

GGTGTCCACA GCCTGCTGCC TGGCCTGGAG CAAATACCTT TGTTAAGTGC TCAGAGGGTA TGGCCCCTCA AATCCACCT
GCAGCTTCCT GGCTGCAAAT ACACTCACTC CATCTTTCA ACTCGCTCCC TGGACCCCTG GTTAACAAT CACTCAATTGCA TACAAAAGGC AACTNGNCAC CANATGGGCA TCCTTGAGCC ATGGTAAACA
CTGAATTTNA GGCTCA

SEO ID NO:1680: (Length of Sequence = 438 Nucleotides)

TACCAGIAGI TCCTTTCCCG CITTATTITI TAGCIGCTIT TIGGGITTIA TACAATGAAC ATGIATTAAT TGIAGAAGAA
AACGATGICA TCCTTTATGA TAAAATCCAT TICCATTITA GCTTTTTTAA AAAAACAAAA AGCIGTIGIG GACAGATGAA
CATCCAAGIA CIGGGCACAC CICCAGCCCT CCCTCTTCCA CIGAAGGCCA TIGCCTATTC CIAGAAAGIT CITTCCCAGG
TATGCAGCTT TCAGITTCCA CITCAGAGGC CACAGIGICT GGGGGAACGG ACIGCCCCCA ATACTAAAGG GAGTCAAAAT
CICTTTAATT NCCGCACTTC CICAGTACCA ACAAGGAAGT CCCTTCTTTA GGGCCACTGG ATGGGAACCT NGGGACCCCC
CITTTTIGAT TGGCAAGCAT TGGGCNTCCT AGGGCCTT

SEO ID NO:1681: (Length of Sequence = 370 Nucleotides)

SEO ID NO:1682: (Length of Sequence = 397 Nucleotides)

ATGIAATCCG CTGCACCAAA CACACCTCA CCAACCACA GGITTITAAG TITGACTGCA CAAACACACT CAATGACCAG
ACCTTGGAGA ATGINACAGI GCAGATGGAG CCCACTGAGG CCTATNAGGI GCTCTGTTAC GTGCCTGCCC GGAGCCTGCC
CTACAACCAG CCCGGGACCI GCTACACCAC GGTGGCACTG CCCAAAGAAG ACCCCACAGC TGTGGCCTGC ACATTCAGCT
GCATGATGAA GTTCACTGTC AAGGACTGTG ATCCCACCAC TGGGGAGACT GATGACGGAG GCTATGAGGA TGAGTATGIN
CTGGGAAGAT CTTGGAAGTT TACTTGTAGC TTGTTCACAT TCCAAAAGGT TCATGGAAAC TGAACTTCGA GCAGCCT

SEO ID NO:1683: (Length of Sequence = 396 Nucleotides)

GECTIGOGCAG AGGAGCOGCT CTOGCOGCOG CCACCTOGGC TIGGGAGCOCA CGAGGCTIGCC GCATCOTGCC CTOGGAACAA
TIGGGACTOGG CGCGCGAGGT GCTTIGGGCCG CGCTGCTCCT GGGGACGCTG CAGGTIGCTAG CGCTGCTGGG GGCCGCCCAT
GAAAGCGCAN CATGGCGGCA TCTGCAAACA TAGAGAATTC TIGGGCTTCCA CACAACTCCA GTGCTAACTC AACAGAGACT
CTCCAACATG TIGCCTTCTGA CCATACAAAT GAAACTTCCA ACAGTACTNT NAAACCACCA ACTTCANGTT GCCTCAGACT
CCAAGTNATA CAAACGGTCA CCACCATGGN AAACCTTACA AGCGGGCATT TTAATTNCAA ACANCAACCA GGGGAT

SEQ ID NO:1684: (Length of Sequence = 417 Nucleotides)

ATCCAGGGA GATGCATGTG GAAATGTGGT CCTCTGGGGT CAGACCCCTG CACGGGACAT CTTGCCTTTN AGTGTGCAGA
GTACATGGGG AAGGGGCTGG GGGCACCACT GTGTACCTGG GCCCAGTAAG GCATTTGCCG TGATTCCCAC AACGGGGTCA
AAAGCTGGCC TTCAGGGTGA CCTAACACCA CCTCATGCCC TGCTATAGAC CTTCACAAAC GACTTCCACT GCTGAAGCCT
GTAGGCTCTG TTTAGAGACA AGAAGATGGC TGGTAATTTA AGCACCGATT TCCCAAGTGC CCACTCTCCT TTGTGCTCTG
TTGGCTTTTG GCCTAAAGCT TNNCCCAGAG TTAGGGTGTA GGATGTCTGT GGTCTGTGAG ATGCCTTTCC CTTCCCCCCT
CTGCTTCAAC CGTGGTT

SEO ID NO:1685: (Length of Sequence = 429 Nucleotides)

 TGAGAGAATA AATGGGTAAT GGGAGGAGAA CTATTTTAAC AAGGGTCCTG GGTTTCTCTT TGCAAACACA GTAGGCTTAA ACTTTGCCTG CTTTTTAAAA TGGCATTTT

SEO ID NO:1686: (Length of Sequence = 445 Nucleotides)

TGICTICATA ATATAACAAC ACTAATACAC TAATAGTAAG ATTAAGTTAG GCAGTCTTCT ACCAAATGTG TAATGGAGAT
TGCCTCAAAA TTGIGTCCAC ATAATCCACG CTCATCTTGC AAAGCGCTAT TTCAGGCACT TTTTTTTGAG AAAGAGTCTC
ATTCTGTCGC CCAGGCTGGA GTGCAGTGGC GCAATCTTGG CTCACAGTAA CCTCTGCCTC CCGGGTTCAA GCGATTCCCC
CGCCTCAGCC TCCCGACTAG CTGGGACCAC AGGCACGNAC CACCACGNCC GGCTCACCTT TGTATTTTTA AGTAGAGATG
GGGGCCTCAC CATATTGGGT CAGGCTGGGT CTTCAATCTN CCTGGACCTC ATGNTCCACC CGCCTTGGGC CTNCCAAAAG
TGCTTGGGGA TTANAGGGAA TNGGGCCACC GGGCTTGGG CCAAT

SEO ID NO:1687: (Length of Sequence = 170 Nucleotides)

AAAAACCAAA TAAAGCAATA ACTTTAAAGA CCTCAGACAC ACACAGTATA AACACCTGGG TAAGGTTTIN TCCGTGTCCA TGTTGACACC GGAACTACCG TTAAAGTGCA AGTTTTGTTT TGTGTTCCTT TGTGCAGTTT CACTCACATG TAAACAAGTC ACTTGGCTAT

SEQ ID NO:1688: (Length of Sequence = 386 Nucleotides)

AATGTGATTT GATGTTAACA CTAGAGAATG ATGACTGTAG AACATTTGAG CAAGTAAAAT AGTAAAGCAC ATAGTGAGTG
TATGTCCATC TAACTGGTAC ATTGATAATT TAGTTTGGGC ACATAAAAGG AATATTTATA TGGCTTCCCA AATGCAGAGT
TACATCTTAT TCGTGTATTT CTCTGAGTAT TTATATCCCG TCTCCTTTTT TCATTCTTAA AAATAAATGA ATTTTCACTG
TTGGCACATA TGAGGCTTAA ATATAAGGAG CATAACACTT GCATTCTAAT TTTTGCATAT ATTGTAAATG TGTCTGGTAT
TTACAGCAAA ATACTGTGTA TCCTTTATGG GTAAACAAAG TGACATTGCA TGCATGTAAT GTGATG

SEO ID NO:1689: (Length of Sequence = 400 Nucleotides)

CITCIGICGG ATCAGCGIAT TCCIAGATIA GGAATICAAA TTAATGAAAA TTCACATATG AAAGGAAAAT CCATTGCTAT
TTCIGGAGAG GACCICAGIC CIGGGCTITI CCCIGGCAIT GCIACCIGGG TGGGIGCICA CCACICAGGI GCIGGIGITG
GAAGGCAGGA GGAGGAACCI GAAATCCIGC CGATIAAGGC TAATIAACAG GGITITAGGIG CCIAATIATC ATGACICAGC
CCGGGACITA TGGITAGCCG TGCAGGCCAG GIGAGICICI TATGGACITC CICTCAGACI GCICTITCIC ATTITGICCI
GATGAGATAT TGACAGICAT GICCACCCGC TICCTCATCC ATTICCCGIC TITGGGCCCI GGGAAGIACG GGGGCCICIG

SEO ID NO:1690: (Length of Sequence = 337 Nucleotides)

AGINATATAC CITTAAAAGT AACTAATGCA ACTGCCAAAN AGGGACAGTG TCAATATCAT TGINITCATT AGAAGGACGG
CTGCCCCACA CTGINAGAAC ACTGCTGTTC CTAACAGTAG TTTACTTTNA GAGGGATGIN AGAATTAGTT TNACCTTAAT
TCCAGATGTG CATGCCTCAA AAGAAAAATC CCATTCTCCT TCCTTTTGGG GAGCACTTTT GGTGGCACCA AGGCTGGTGT
GGGGTAGTGG AGAGAGCACT GAGCTTAGAG TCACAACCAG ATGAAACTGC TCTGGTCTTC ACTAGCTGTG TGACTTGGGC
AAGCAGCTTG CAGTCTC

SEO ID NO:1691: (Length of Sequence = 372 Nucleotides)

TCATTCTCCC AAAGTGCTGG GATTATAGGC GTGAGCACGT GCGCCCAGCC TTACTTATTT TTAAATCAGA TTTTTTÄATC
AACTAAAACA GCTATGAGTT AAGTACCTGC CCTGCAAAAA TTTTTTTGAAA AAGTTTTTGG ATTATGAAACA GTTATGAACAGT CTAAAAATTTA TCTGATACTT CTCTAACAAG TGAGTGATCT CATGTAACCC CAGTTTGTAT

CTTAAAGGCT GCAGCATAGA ATTGAGCTGT ATAACAGTGT TAGAACTGTC AAGTGATAAT CACAGAACAG TTTGTATCGG TTTTATAATT CTCATGTCTT GATCAGATCT GAAGGGAATA GGCATACCCT CC

SEQ ID NO:1692: (Length of Sequence = 360 Nucleotides)

TITTITIGC AAAAATAGTA TATATTATT ATGIACAACA TGIATTIGA GATATGIATA CATTGIGGAA TGICTAAATT GAGGCTAACAA ATACATTATC TCACATACCA TGITTITTIG TGGIGACAAC ATCIAACAAT ATAGACCATT TCACAAATTT GCATGITATC TITGIGCAGG GGCTATGCCA ATCITCICIG TATTTTINCA ATCITGGIGT ATGIGCIGCT GAAGCACACA CCCTAATTCC TITCATTITA GGMICTAGIT AACCITICIC TTAAGTATAA CCATGTATTT TGITAAGCAA TATCITTITTA TTACAAAAAAT GCCATTTITTT TCIGGMTAGG AAAATTGATT

SEQ ID NO:1693: (Length of Sequence = 378 Nucleotides)

GACAAAAAGA GGGGICIGGC TGCCGATGIG GAAATTIGIT TTGIGGACTI CACCGITACT CIGACAAGCA CAACTGICCG
TATGATTACA AAGCAGAAGC TGCAGCAAAA ATCAGAAAAG AGAATCCAGT TGITGIGGCT GAAAAAAATTC AGAGAATATA
AATTACTICT TGIGAAGAGA CIGAAACTIT GITTITATIT TAATATATCG TAGGAAAACA TTAAAGAGCA GATGCATGGC
CATTITUCTT TGATGITCTC CAGAGTITTA CATTACACTI GICTGICTTA TAATTGATAT TITTAGGGATG TTTGGGTGTT
TGTTACAGGC AGAATTGGAT AGATACAGCC CTACAAATGT ATATGCCCTC CCCTGAAA

SEO ID NO:1694: (Length of Sequence = 362 Nucleotides)

AATGCACTIT ATTGCTCCC AGGGAGTGGG ATGCAGGATC AGAGTGGACA CGCGCAGGGG GCTGGTGTGG GGAGCAAAGC NCCGGGCCTG CCCCGGACCC TGGTTTCCCT GAGGACCAAC GTGAATGGGG GCCCCACTGG AAAGATGCTT GGGGCTGCAG AGGGGGGGAGAAGC AGGTTGCTGG GTGGTGCCCT CAGCTCCTGG CAGGGTTGAC GGGTGGTGCC CGCTGGGCTC TGCCAGCCGA TGGTCCNCTG GCACCTGATC CTGTCTTCCA GCTTCACTTC CGGGCCTGCT CGTAGTTGTC AGTGAACCAA GCACAGGTCT CCTTGACCGA CTGCTTTNAA GGGTGTGAAN CG

SEO ID NO:1695: (Length of Sequence = 411 Nucleotides)

TTAATACAAG GGGTTTGAAC TGGACATCCT AATGATGCAA TTACGTCATC ACCCAGCTGA TTCCGGGTGG TTGGCAAACT CATCGTGTCT GTCCTGAGAG GCTCCACAAT GCCCACCCGC ATCGCCATTC TGTAGTCTTC AGGGTCAGCT GTTGATAAAG GGGCAGGCTT GCGTTATTGG CCTAGATTTT GCTGCAGATT AAATCCTTTG AGGATTCTCT TCTCTTTTAC CATTTTNCTG CGTGCCTCCACCC CTTGTCTCT CTCTCTCTAG CTTTTTAATT CATGAATATT TTCGTGTCTG TCTCTCTCTC TCTCTGTGTT TCCTCCCAGCC CTTGTCTCG AGACGGTGTT TTCCTCCCTT GCCCATTATC TTTTCAACTC CCAGGGCTAC CCATTTCAAT GGTGGGTCGT T

SEO ID NO:1696: (Length of Sequence = 280 Nucleotides)

CTITIGICATE TITITACECT TIACAAAAAG CAGATITEGI ATTCAGAAAA GCCTGCAAAT ACAACATTGC TIAAGAGAAC CTIGIAAACAC GITTGGAATA CAATGCAACA CAAGTCAGCA AGGACAGGGG TAGGTCCAAA GGAGCCAGCT AGGGGGAAAG GTGACAGAAA AGGAGAGGGA AGGATGGAGA CAGACATCAC CTGTGGTCTC TAAGGGGGGCC NTGTGTTAA TITATAAGGT TINCINCCCA CAGGAGTTCT NNTGTGATCT ATCCGTTCAT

SEO ID NO:1697: (Length of Sequence = 418 Nucleotides)

 GTACCCCAGA GCTTNGTGGG TGAGTATTCC ACCTGCTTAC ACACCACTGA AGCCACAGCC AGCCAGTAAC TAAGGGGCAA GAAAGAGCAT TGTCCAAGCT GGCTCTTING GGGGGTCCCC CATTNGGCCA CAAAGGCCTC ACCCCCACC CCATCCCCGT AACCAGAAAC CACCTTGA

SEO ID NO:1698: (Length of Sequence = 376 Nucleotides)

ATTITIATIG TITATITACI TATTITITAC CCITITITCA AGAGATGGGG TCTCACAGTG TIGCCCAGGC TGGACTIGAA CTCCCACTCC TGGGCTCCAG CAGTCCTCCT GCCTCACCTT TCCAAGTAGC TGGGCTATA AGTACACACC ACCATGCCCA GCAATATITT AATTICTGTA ATGIGTCATT TAGCCAGTGA TTGTTGTATT ATAATAGAAT CACAGAAATG GAGGGACTCC TAGAGGGTAAT CAAATCTGGT GGTTTTTAAG CCTTTTATTC CCTCTAAAGG GATAGTAAAA CCATTAAAAA TATAATTTTT CCCCAATTATG TAAGCCCAGTG AAAGCTGACC TYCTGGTTTA GAGAGGAACA CAGATG

SEO ID NO:1699: (Length of Sequence = 365 Nucleotides)

GGTACATGIG CACAACGING GNGITIGITA CATATGIATA CATATGCCAT GITAGIGIGC TGCACCCATI AACICGICAT TTAGCATIAG GIATATCICC TAATGCTATC CCTCCTCCCT CCCCCIACGC CACAACAGIC CCIGGIGIGT GATGITCCCC TTCCIGIGIC CATGIGITCT CATTATTCAA TTCCCACCTA CGAGIGAGAA CATGCIGIGT TTGGITTITT GICCITGCGA TAGGCCAGATG CAGCIACTCT TAATGIGCAT ATTTTCATCC TAGAACATIG GAGAGITCCT GIAAAAGCCT TGIGITCCAG GAGAGAAGGAG ATCCIGACCC TTCTGCTGAT GGCAGCAGTC AGGGG

SEO ID NO:1700: (Length of Sequence = 397 Nucleotides)

AAAGGCAGTC AAGCAGGAGT TAAACAATAT GGACCTAACT CTCCTTATAT GAGAACATTA TTAAATTCCA TTGCTCATGG
AAATAGACTT ATTTCTTATG ATTGGGAAAT TCTGGCTAAA TCTTCCCTTT CACCCTCTCA GTATCTCCAG TTTAAAACCT
GGTGGATTGA TGGGGTACAA GAACAGGTAC GAAAAAATCA GGCTACTAAT CCTGTTGCTT ATATAGATGA AGACCAATTG
CTAGGAAGAG GTCCAAACTG GGACACTATT AACCAACAAT CAGTAATGAA AATGAGGCTA TTGAACAACT ATAAGAGTA
TTTGCCTCAG GGGCCTGGGA AAACATTCAG GACCCAGGGA ACCTCATGCC CTTCTTTTAG GTTCAATCAG ACAACGT

SEQ ID NO:1701: (Length of Sequence = 245 Nucleotides)

GTCTAGGAGG AGGCCTTCTG CACAGAGCCC CTGAAGAACA CAGGCAGAGG CCCCCCACTT GGCTTCTACC ACGTCCAGAA CATCGCAGTG GAGGTGACCA AGTCCTTCAT TGAGTACATC AAGAGCCAGC CCATTGTTTT CNAGGTCTTT GGCCACTACC AGCAGCACCC GTTCCCGNCC CTCTGCAAGG ACGTGCTCAG CCCCCTNAGG CCCTCGCGCC GTCACTTCCC TCGGGTCATG CCACT

SEQ ID NO:1702: (Length of Sequence = 349 Nucleotides)

ATCTGTTGTC AGCACAGITT TATTTGCTGT GGAATCCATG AGAGCCGGAA GCATCGTTGG GGCCGTGGCT AGCAGAGCTC
ATGGTGACCA GTCCTGGGCC TGACCAATGG GTGATTACAT TTAAAAACCA AAACAAAAACA AAACAAAATA CCAAGAACAG
ATCACTTGCC ATGGACATCA GTAATCTATT GGTAATGGTG AAAATTTCAT GAAAATTTCC CCTAAACCAT AACAAAAACT
GTCCTCCTTA CCCCAAAAGT GCTGGAGGGA AAGATGGTTG CATGGCTTTG ACCTCTCTTT GAACTTGAAA TGCTACCTTC
CTACCCGGAA AATGCGGCAC ACTATACTT

SEO ID NO:1703: (Length of Sequence = 419 Nucleotides)

GAGCCCCIGC CCTCCAGAAG CTCACATCCT CCTACTCATG GCAGACAAAT PARCETCAT TACACTCAGGCAGGCTAAGT GTGGCAGCAG ATGTAGTATG CAGTGCAGAG GTGGCCATGG TTGCNAGGGC AAGGAGGGCT TCCTAGCATG GGCGTTATTT GACCAGAGGC TGGCGGTGGC TTTTGCTAGC AGTGTGATTG TNATCTGAGC CAGGGACAGA TACCTCINTG AGCCTTGGTT TCCTCATCTG TAAAGTGGTT AAAGACTGAN TAAAGCAAAA TATGTGCAAA CAGTCTGTGA ATGGGGAAGT AACAGATGTT GCTTTCTATT ATGTTCTCTC CTAGCCATGA ATATCAATTA TTTCAGAAAT GAAAAGGGAT CCTGCACCCA ATTTCAAATC AAGCAAGTTC ACCTAGAGG

SEO ID NO:1704: (Length of Sequence = 372 Nucleotides)

GCTTCCCGAA GGTCTTGGAC GAGCGCTCTA GCTCTGTGGG AAGGTTTTGG GCTCTCTGGC TCGGATTTTG CAATTTCTCC
CTGGGGACTG CCGTGGAGCC GCATCCACTG TGGATTATAA TTGCAACATG ACGCTGGAAG AGCTCGTGGC GTGCGACAAC
GCGGGGGAGA AGATGCAGAC GGTGACCGCC GCGGTGGAGG AGCTTTTGGT GGCCGCTCAG CGCCAGGNTC GCCTCACAGT
GGGGGTGTAC GAGTCGGCCA AGTTGATGAA TGTGGACCCA GACAGCGTGG TCCTCTGCCT CTTGGCCATT AACGAGGAGG
AGGAGGATGA CATCGCCCTG CAAATCCACT TCAACGTTCA TCCCAGTCCT TC

SEO ID NO:1705: (Length of Sequence = 426 Nucleotides)

GATGCCTTAT TRAGICCATT TEGTGAGGIA ATGITITCTI GGATGTCCTT GATGCTTGTA GACATTTGTT GATACCTGGG
CATTAAAGNG TRAGGTATTT ATTCCAGTCT TCACAGTATA GGCTTGTTT TAGCCATCCT TTTTGAGAGG ACTITCCAAG
AATTCAAAAG GGATTGAGTG TTGTGACCTA AGCCTATGGT CACTGCAGCC ATTTCAGCAC TAGAGAGTGC CCTAAGCCCC
GGAATGCTGC AACTCTTACA GACTCCTTGA TACACAGCTT TGGTAGATTT TGGGAAAATA AGGGAGAATT CCCTGGGGTT
ACCAGGTAAA AAGTCTCTCC CACTTCCCTC TCTTTCTGGC AAAGGAAGTC AGTCTCTGCA CCAGGCTGCC TGGAGTTTGG
GGGAGGGATA AGGCGGTCAC TCTAAT

SEQ ID NO:1706: (Length of Sequence = 412 Nucleotides)

ATTITATITC CITACATCGA AGAAAATGIT AAAGAGTATC TACAGACACA TIGGGAAGAA GAGGAGIGCC AGCAGGATGI CAGTCTITTIG AGGAAACAGG CIGAAGAGGA CGCCCACCIG GATGGGGCIG TICCTATCCC TGCAGCATCI GGGAATGGAG TGGATGATCI GCAACAGATG ATCCAGGCCG TGGTAGATAA TGTGTGCTGG CAGATGTCCC TGCATCGAAA GACCACTGCA CTCAAACAGC TGCAGGGCCA CATGTGGAGG GCGCATTCA CAGCTGGGCG CATGAAAGCA GAGTTCTTTG CAGATGTAGT TCCAGGAGTC AGGAAGTGGA GAGAGGCCCG AATNAAGGTG TACATCTATT CCTCAGGGAG TGTTGAGGCA CAGAAACTGT TATTCGGGCA TT

SEO ID NO:1707: (Length of Sequence = 434 Nucleotides)

GEGETETCEC AAAAAAAAA AAGATTCIAG GCATGGEGT GEGITGACIG TAGITCCAGC TACTCCAGAG GCTGAGGIGG
GAGGATIGGI TGAGCCIGGG TGGATGAGGC TGCAGTGACC CATGATCATG CATGGGAGAC AGAGCAAGAC CTTGICTCAA
GAAAGGAAAG AAATCACIGG CTCTTCIGIA AAAAATGATC TGTTAAGAGI AATTGAAAAA ATAAATACAA GIAATAAAAT
AATCITTCAT TTAAGAAATA CTACCAAAAT TAACATGGAG ATCIAGCAAA AAGTCAAAAG CAGCINGGCG TGGIGGCTCA
CACCIGIAAT CCCTACACCT TGGGGAGGCT GAGGCGGGAG GNICGCCIGA GGICAGGAGT TCGAGACCAG CCTGGCCAAC
AGAGCCAAGT CTCTACTTAA ATACAGATTA GCTT

SEO ID NO:1708: (Length of Sequence = 440 Nucleotides)

GGACCAGGAC TCCAGCACCT TCCCTGGCTG CATCAACAAT GCCACACTCT TTCAAGATGA GATAAACTGG CGCCTCAAGG
AGGGACTGGT GGAAGGCGAG GATTATGTGC TGCTCCCAGC AGGTGCTTGG CATTACCTGG TCAGCTGGTA TGGTCTAGAG
CATGGCCAGC CACCCATTGA ACGCAAGGTC ATAGAGCTGC CCAACATTCA GAAAGGCTCAGGTAACTGCT
GCTTGTCCGG CACAATGATT TGGGCAAATC TCACACTGTT CAGTTCAGCC ATACCGATTC TATTGGCCTA GTTATGGCCA
CAGCTCGGGA GCGGTTTCTG GTGGAGCCCC AGGAAGACAC TCGGCTTTGG GCCAAGAACT CAGAAGGCTC TTTGGATAGG
TTCGTATGAC ACACACATCA CGGTTCTCGA TGCGGCCCTT

SEO ID NO:1709: (Length of Sequence = 404 Nucleotides)

TTIGICITAT GIAGAATIGC CIATAGIAAG AAAACCCAGI AGAGAAAGIG GITTINAGAC CATTCGGCAG CIGCITIGGA
CACCIGGAGC CATITCITIT ACAGATGAAG ATGCATIGIG TCATTGICIC AGGATCCICG TCCIGITGCI TCICIGGCCA
CAAATIGITC TITACCAAAG ATGATTITAT TICACIGICT TIGAAAATCA TICITITATAG GIAGAATATG AAGATTCICT
GAAATGATIC CAAAATGCCA AACTCAAACA CIATIGICCG ATTTCTTTAC TIGCAACAAG AGAGTAGGAAG GGACAGTATT
TGITTIGIGA TGITGGGGCG TTCATCAGGG AGAGAATTIG AGATAAGIAG GAATAGCAAA TAGGAATAGI GAAATAACCT
AGAT

SEO ID NO:1710: (Length of Sequence = 187 Nucleotides)

GGTGATCTGC CGACCAGAGG CCTTAAACTC TGGTGTTGAG TACTACTGGG ACCAGCTGAA CGAGACGGTC TTCACTGTCC ATTCCAACAG CAGGAGCAGC GAGGGSCTGG ACCAGGCAGA GCACATGGAG GACAGCAGAG ACATGGGCTG ATGAATGCAT TGGGCTTCAG CCGACCTGCA CTCAGTG

SEO ID NO:1711: (Length of Sequence = 313 Nucleotides)

AGGGCATGI NATCATITNA ATGATGINAT CITTGGTGIT TCCCTCATTA GCTGTAGACT ATCCCCTCTC CTCCCACCAC
AATGITTCTA TGATGAGITA CAAACAGAAA GGAAATCACA TTTTCATACT AAAAACAAAA TGATCAGAGC CITGATTTCT
CCACTAGAAA CTACACGTAC AGTTAAGAGT CCACATGCAA CACCTTAAAT CACAGACTGA GGACCTCACA TTCTGACCTG
GGAGTCTCCT CCCCTTCCCC AGCCTTGGGC TAGCTTTGGC CTAGGCTCAG GTAATACTGA CACCCACAGG CGT

SEO ID NO:1712: (Length of Sequence = 202 Nucleotides)

TTITGGIGGI TICCICITIA TIGIGIGCCI CCIACCITCC CCCACAATTI CAGICCCITC CAACACCCCA AAAAGAAGGA GIGAAAGGAA GGGATTGCIG GGGITCIGAG CCCTIGGCAG TCAGAAGGAC AGAACCAAAC ATCACTGGAT GIGACACAGC TGCATCAAGA AGICTACAGC AGTATGGGAA GCGGCAGAGA AG

SEQ ID NO:1713: (Length of Sequence = 253 Nucleotides)

TGATTCANTG GGTCTGGGAT AGAGTCTGGT ATTCTGCATT TCTGACTAGC CTCCAGGTGA TACTGATTCT CCTCATCTAG
GGACCTCGCT TTGAGTAGCA AGTGTTTAGG CCACTTACTA GCAGGAACTA AGCACAGTAT CCTACAACAG CAAATGTCTT
TCCAACAAGA AAGACGAGAG CAAATNCTGA TGCCACATCT GCACTGCCTC AGAAAATAAA GAAGGGATGA GGAGCCCCCC
AGTGGCACTC TGT

SEQ ID NO:1714: (Length of Sequence = 299 Nucleotides)

GGTGCAGCTG CTTTGAAAAA TGACTTGGCA GCACCTCAAA ATGTTAAACA GAGTTACCAC ATGACCCAGT AATTTCACAC
TTAAGGATAT ACTCAAGAGA AATGAAAACT AAAAACATAC GGCTACCCAA AAACTTACAT AAGANTGTC ACAGCAACAT
TATTCATAAT AACCAAAATA TGGAAACAAC CACAATGTCC ATCAATTGAT AAMTGGGTAA AGTCTGGCAA ACTCACAGRA
TGGRATATTA TTTGGTGGTA AAAAGGAGTA AAGAACTSNT ATGTACTACA ACATGGGTG

SEO ID NO:1715: (Length of Sequence = 371 Nucleotides)

TTTTTTTAC COGGGGGTTC CTGAGTTTAT TTGGGGCACA CCCGGACGAG GGCCCTGCAC CTAGAAGAAG GTGTTGGGCC
*CTTGGTGGT GAAGCGTGGC TTGTGCTGAC GGCCCAGGC CCGGTGGGC AGCGGGAALT TCATCTTGACAG CCGGCGGGGC CTCGGGCGCG
TGCTTGACAG CCGGCCGGCG GCACTTGCTG GCCGCGATCT CCTCCACCTT CATGATCTGA ATGGAGTGGG CTCGGGCGCG

GTGCCGGGCA CCCATGTCTC GGTAGCACTG GGTGACAGCG CCTGCGGTGG TCAAGTCCCG GTATTCCCGG TACATGTTGT GGGTGCCGCT CCGGGAGTCA TAGCGCAGCA AGATCCCGAA GTTCTTCAAC C

SEQ ID NO:1716: (Length of Sequence = 265 Nucleotides)

GTGCAGAATC TGCTCCTGGA CACCCACAGG GGGCTGCTGT ATGCGGCCTC ANANTCGGGC GTAGTCCAGG NGCCCATGGC CAACTGCAGC CTGTACAGGA GCTGTGGGGA CTGCCTCCTC GCCCGGAACC CCTACTGTGC TTKGAGCGGC TCCAGCTGCA AGCACGTCAG CCTCAGCTG CCACCAGGGC GTGGATCCAG GACATTGAGG GAGCCAGCGN CAAGGACCTT TNCAGCGCGT CTTCGGTTGT TTCCC

SEO ID NO:1717: (Length of Sequence = 350 Nucleotides)

CAGCCCCCCC AGCCCTCTGG CCCCCTCCAT CTCTTGTCCG TTCCCACCCA CCCCCCTCCT CGGCCCGAGC CTTTTCCCGG
TGGGTGTCAG GNTCACTCCC ACTAGGGACT CTGCGCTAAT TACCTGAGCG ACCAGGACTA CATTTCCCAA GAGGCTCTGC
TCCAGGAGTC CAGGAAAGAC GAGGCCACCTT GGCCGCGGG CCTGCTGGGA CTTGTAGTTG CCTAGACAGG GCACCACCCT
GCACTTCCGG ACCCGCGCTG GAGGCGCCGT GAGGTTTGGT GTCTCGAAGC AGCAATTAAA AAGCAAGAGG ACTTCATGAC
CACCATGGAC GSCAATTAGG AGAAGATCAA

SEO ID NO:1718: (Length of Sequence = 379 Nucleotides)

GACATGGAGA CTCACATGGC TGCAGAACAC TGTCAGGTGA CCTGCAAATG TAACAAGAAG TTGGAGAAGA GGCTGTTAAA
GAAGCATGAG GAGACTGAGT GCCCTTTGCG GCTTGCTGTC TGCCAGCACT GTGATTTAGA ACTTTCCATT CTCAAACTGA
AGGAACATGA AGATTATTGT GGTGCCCGGA CGGAACTATG TGGCAACTGT GGTCGCAATG TCCTTGTGAA AGATCTGAAG
ACTCACCCTG AAGTTTGTGG GAGAGAGGGG GAGGAAAAGA GAAATGAGGT TGCCATACCT CCTAATGCAT ATGGATGAAT
CTTMGGGTCA GGATGGAATC TGGATTGCAT CCCAACTCCT CAGACAAATT GAGGGCTCT

SEO ID NO:1719: (Length of Sequence = 197 Nucleotides)

CCTATATTIG TITAATITAT TITAAGACCAC CTCCTTACAA CTTCCAGAGA GAAAATACAA AACAAGAAAC AGACTIGGIT TCAAATGCAT AACCAGGIGC TGGAGITTAA AGCATTACTG ATAACATTGI TACAGAAGAA TGGCAGCTTA CTCCAGGGCA CTTCAGTATT CCTGAGGAAT AAACATGATT TCGGAAG

SEO ID NO:1720: (Length of Sequence = 203 Nucleotides)

GAGGGCGGG CAGAGGAGC ATGACGGGA GAGTGAGGAG GAAAGAGGAA AGGAAGGCCA GGGTGGGAGG AAGGATCANC
TAAATCTGAG GGAAGAAGAA GGAAAGGAGA GGCCTATTT CATAGCAGAT GCAAATRAAG GGNCTTGGGG CTAKTCAGGA
AGAAAGGGAA AGGGAAGGAA GGCAAGAGGA AGGGGTGAAG GGA

SEQ ID NO:1721: (Length of Sequence = 326 Nucleotides)

GGTGCAGCGA TGTTTAATGG CAATTCGTAT AAACCAAGCC CATGCACAAG TAGAAAGTGC CCGTGGAGCC GGCAGGAGGC
CCCCGCCGCG NTAGAGAACC ACAAGCCCGG CCGTGCAGCC CTCCCCGCGG CGCCTTAAAT AGATTCTTCA CTATACTCTG
TATGTTACAG TATGTACAAG ACCCCTCCCC TCGGGGGACG GGGCGGACTN CGCAACGNGT TCCTATGTAC ACCACCTCCC
CTTTCGGCCC TGAGGTCAGT GGCCAGAGTC GGGTGATGGG GTAAGANAGG GCCAGAGAGG GAGGAAACAG ACGCAAACAT .
GCGGAG

- 45

SEO ID NO:1722: (Length of Sequence = 291 Nucleotides)

TGITTTAAA AATGAGAAAA TTTGGAGAGA GAATACTATT ATGTCAACGG TACAAGACTC TGAATCTIGA AGATGTAGAT
GGATATAAATA TTTAGACTIT ATATACACCC ATAGATATGI ATTTATATAT GCATACGTIT TGTATAAATT TACAATTGAC
TTTTTGTATT CTCTTTACTG TCATTACAAG AATGAGATGG AAACCAAAAT AGITGTNCCA TCCTCTTACC CAAAGAGGGA
TACTGAAAAG TCCGGTATGI GCATGCACTT GITTCTCTGG GGTCAAATCT G

SEO ID NO:1723: (Length of Sequence = 369 Nucleotides)

GATTGCCCGC TCCCTCGATT CCTTCCTGTT GTCTCAGAA GCTGCTGTG GCTTGCTAAA AGGGACAGCA CTTGTCCTAG
CCCGATTACC TTTGGATAAG ATTACCGAAT GTCTTAGTGA ACTATGTTCT GTTCAGGTTA TGGCATTGAA AAAGCTGTTG
TCTCAAGAGC CCAGCAATGG CATATCCTCA GATCCCACAG TGTTCTTAGA TCGCCTTGCA GTGATATTTA GGCATACCAA
TCCCATTGTG GAAAATGGAC AGACTCATCC GTGTCAGAAA GTCATACAGG AAATAINGCC AGTTTTTATC CGAGGACTCT
AAAATAAGCA CCGAGCTGNA TAATCGGATT GTAGAGCGTT GTTTGCAGG

SEO ID NO:1724: (Length of Sequence = 231 Nucleotides)

ATGIATIGIT AGITCGATTC CITCAAATIT TATACATATI TACTITCIGI TAAAGAGAAA AGGATAAAAT GGIATAAAAA AAGATAAAGC TATTAATTAA GCACGAGAGA GAAGATAAAT GGATATTITC CCTGTGTGAG GCTAAGACAG AWGCAAATCT CGITANGAAA AATGCCACCC ACACAACAGG AANITTATCC AAAACAAAAC AAAAGCAGIT ATAGANCCCC T

SEO ID NO:1725: (Length of Sequence = 317 Nucleotides)

GTGCAGGGTA GGGTACATAT GGCTCTGTCA GAAGAATACC ATGATTTAAG GGAAGAAAGT ACACAAGGTA CATGGAGGGT
ACACAGGGAA AGTACAT. A TAAACATGGA CGTGTGCAAA TAGGAAAGAC ATGACTCAGC ATGCTAGACA AATTGCACAT
GCCTACCCAA ACACGCT. A GGGCAGACCC ATGACCATGA GAGGGGCACA CGTAGCTGTG AATGCAGGGC ACCCGAGAGC
ACATGTKACT KAACATGAAG AAAGCATACG GGAAAAGCGT GTKTACACAT GNGCATGTTC AGTGGGGCAC ACGCAGG

SEO ID NO:1726: (Length of Sequence = 282 Nucleotides)

CTCTTGAACC AGATGAGCAG CCACCGGAAA CAGAAGCAGA GAGAGCCGGA GTCCTGGGAA TCCAGGAAGT CGCAGAGCAG
GGGGTCCAGC ACCCTCAGGA GCAGCAGCAG TCGCCCGAKT TGCCGCTTCA TGGTCTCCTG GCTCTCTTCA AAGTTCCCTT
GCACGAGCTC CATGAAGCCA CAGAAACACC AGAAAGCATC CACCTCGTTC TGAATGACGT AGAGGATCGG GGAGAGAAGA
TCACTCATGC CCTGGACGTA GCCGAGGTCG AAGTGATACA TT

SEO ID NO:1727: (Length of Sequence = 285 Nucleotides)

GAGIATIGAT TICAGGCAGG ACCCAGGICC CAAAATGITA GAAACAGITA TCCITITICC CICIGAGITC GITATICICI
GGGGCCCCAG TATCCGIGGC TIAACAACCC GGCIGGATAG AAGGCACCIC TITCCCCACG TICCAACAAG ATCCCAGAGC
TGCTTCTCAT TGGCTCGICC CIGAGICAGI CACACTGGAC CGGAAGGIGA AAGGCCCTCA TIGGCCAGNC CCGAGICATG
TGCCCACCCC TGGGGATCCA GCTGTGGGNC INCITITAACA GCATT

SEO ID NO:1728: (Length of Sequence = 394 Nucleotides)

TITITITGAT GAGGAGATAT AGCAAAGGGT CATITGCCCC TCCTTCAGAA AACTITTCTC CAAATCTCCT TIAAACATAC
TGCCTTATCT TTCCCTCCAT AACTCCACCA GTCTCTCCAC ATCCCCCTCC AAATCTCTGT ATACATAGGC AAGAGAGGGC
GATTCCCAGC ACAAGTCTAG TCCTGGGCGA AACTTCCATC TCTTTCCTCG CATACCTCCT GTCTGGGTAT GGGGATAAGG
GAGAGTATGG GATTTTGTTC TCCATTACAT GCTTTTTCAA AATTTCTGTA ATATGTGGCA CTTATAAAAC AACAACAGAC
AAAATGATAT CGGGTAAAAC ATGCAACTGA GAGCAATTTG GGGAAAAATC CTCAGGNCAC AAAATGTATT ACTG

SEO ID NO:1729: (Length of Sequence = 301 Nucleotides)

GGAAGTTAAA GTATTTATTG ATGTGTTAA ACTGTGTACA TTCTCCACAG ATCATATTAA GGNGTTKTA GGKGAAGTTT
AATCTGTGCA TAGTGGGTAG YGACATGAWT AGGGTCAAAG GGGAGGYAAA AGGAAAAAAA CAAAACAAAA ACAGTCACAG
GAAAWTAAAA ATACACCMCA GGTTACCAGA ACCTTCAGGT TTAAAATAAA ANENAAGNAA AAGCAGAAGC AGTGAGCATC
GGCATCAACC TGTACAAGCA TTACAAAAGG CTCCTGTGAC GGAAACACAA TTGTTCAAAG G

SEO ID NO:1730: (Length of Sequence = 312 Nucleotides)

GACGRACGCT CTTGCCACGC CCTGAGCGTG TACACATGAT GINTTCTATG CATTCACCCT GCCCCCAGC CCGCCCTGCA
GAGGACAAGA TGGGTGGCCC CGGCTCCCTT TCCCCTAACC GCCCCTGCCC GCTGTGCAGC CGTGTGCGTT
TCTGTGTCAC TGGCGTGTCA CGTGATGTAG CCGTGTTTGC TGACATGAGC CCCTGCCCCC TTCTCTGTTT CTCCGTTGGT
TTCTAGAGGT CTCTCCCTCC CCTTCTCAGA GGGGACAGGA CTCCTGGGGT CTGGCTCGGG CCCAGAGCCA GG

SEO ID NO:1731: (Length of Sequence = 392 Nucleotides)

ATCGGCTATG GGITCCGGTG CGTGACAGAG GAGTGCCCGC TGGCAGTCAT CGCTGTGGTG GTTCAGTCCA TCGTGGGCTG
CGTCATCGAC TCCTTCATGA TTGGCACCAT CATGGCCAAG ATKGCGCGGC CCAAGAAGCG GGCGCAGACG TTGCTGTTCA
GCCACCACGC GGTTATTTCG GTGCGCGACG GCAAGCTCTG CCTCATGTGG CGCGTGGGCA ACCTGCGCAA GAGCCACATT
GTGGAGGCCC ACGTGCGGGC CCAGCTCATC AAGCCCTACA TGACCCAGGA GGGCGAGTAC CTKNCCCTGG ACCAGCGGA
CCTCAACGTG GGCTATGACA TCGGCCTTGA CCGCATCTTC CTGGTGTCGC CCATCATCAT TTTNCACGAG AT

SEO ID NO:1732: (Length of Sequence = 352 Nucleotides)

SEO ID NO:1733: (Length of Sequence = 321 Nucleotides)

TTTTTTGTT GITTGTTGT TTGTTTGCAG AGICTTGCTC TTGATCTATC TCCCAGGCTG AAGIACAGTA GTGTGATCTC
GGCTTGCTGC ACCCTCTACC TCCCAGGTTC AAGCAATTCT CATACCTCAG CCTCCTGAGT AGCTAGAACC ATAGGCACAC
GCCACCATAC CTGCTAACTT TNCTATTTTT AGCAGAGACT GGATTTTGCC ATGTTGGCCA GGCTGGTCTC GAACTCCTGG
CCGCAACTGG ATCTGCCCAA CTCAGCCTTC CAAAGTGCTG GGATTACAGG CATAAGCCAT TCATGTGCGG TTKTTCAACT
G

SEQ ID NO:1734: (Length of Sequence = 208 Nucleotides)

AAGTCAACGT ATCTATTTTT ATTATGAAAC ATTAAATTTT GACACATTGC CTCATTTGCT TTTTTAAAAT CTATTATCTG
ACTTAAACCT ATTCAGCAAA AATGCCAATA AATTATATTA ATCATACTTT GGGTCTTTTT AAAACTAGGA ACATAATATG
TTTTATGATA AACAATAATA CTAAATCTGA GTTGTATGAA CTGTTAAC

SEO ID NO:1735: (Length of Sequence = 347 Nucleotides)

TCTATTACCT GTACACTATG GTTTATACGT TGGTGAGTTT CTAAGGGGGA AGCCCCCAG GGAGGGGGCAC TGGTGCCCCC CCCACCCCGC CTCGGGGGG GAGGGGGCAC TGGTGCCCCC AGCCTCTCCA ACCCCCAAAC TGCTGCTGCG GGGAACCCCC CCCACCCCGC CTTCAGAGCC CTCCCCCTTG GACTAGAGCG

GCTGGGCAGA GCTCTAAACA GGGGCAGGGG CTCCTCTGCC AGCCTGTGGG CATGGCAGTC ATTCCTGGAA GGGGCAGGAC CTCCGGCCTT GTCCATTTCG GGGGGAA

SEO ID NO:1736: (Length of Sequence = 356 Nucleotides)

GACACAGGA GGGGAACAAC ACACACTGGG GCCTGTTGGG GAATGGGGGG TGAGGGGAGG GAGAGCATCA GGACAAATAG CTAATGCATG TGGGGCTTAA AACCTAGATG ATGGGCTGGG CGTGGTGGTT CACGCCTATA ATCCCAGCAC TTTGGGAGGG TGAGGCGGGC GGWTCACGAG GTCAGGAGAT CAAGACCATC CTGGCCAACA TGGTGAAACC CCGCCTCTAC TAAAAATACA AAAATTAGCC AGGCATGGTG GTGCGTGCGT GTAATCCCAG CTACTCAAGA GCCTNAGGCA GGAGAATCAC GTGAACCTGG GAATCGGAGG TTGCAGTGAG CCAAGATCAT GCCACT

SEO ID NO:1737: (Length of Sequence = 324 Nucleotides)

TGITTICTAA TGATTITTAA TITTICAGAG GAAAATAATI TCAAGAAATA AAACITAATI CCCCTGAGIC CITATIGAAT
TAAATATIGA AAAACAATGA AIGAATGAIG CATICITATI AAIGGACIGI AAGAAACIGA TATAATGGAC TICATICTAC
AAITCGGITT CITATIGICI TACACATGCI CCTCGAACIT AAACATITTA GGACCITAAC ACCATTICCC TAGIACAATI
ACTAAAAAGAA AGCITTGGAT AATATAATAT CAGGGAAGAT AGTACAACAT AGTGAAGGAT GACATAGGGN AGATGTGAGG
AGCA

SEO ID NO:1738: (Length of Sequence = 316 Mucleotides)

GGCACCCIGG GCATGICCAG CCIGGAGCAG CTGGAGCAGA ACTIGGCAGC AACAGAGGAA GGGCCCCIGG AGCCGGCTGT
CGIGGATGCC TITAATCAAG CCIGGCATIT GGITGCTCAC GAATGICCCA ACTACITCCG CTAGGCCCAT CATGGCTCAG
GCTGCCCCAAG GCTTTINIGI CACCICTITI GITCTCTCAC ACTGACCAGI CTTGGCCTTA AGCTGACTTA GAAGGGTTTT
TCTGAATIGI CTAGATCCAT GCATTATITI TCTAGCTTCC TGCCTTGCTC CCTATTCACT TTACACTGTG AAAGGT

SEO ID NO:1739: (Length of Sequence = 398 Nucleotides)

CAAAAACCAT CTCAGGATAC TGAGAAGCCT CTGGAACCTG TGAGTACTGT TCAGGTAGAG CCTGCAGTTA AGACTGTAAA CCAACAGACT ATGGCAGCAC CAGTAGTCAA AGAAGAAAAA CAACCTGAGA AAGTCATCAG CAAAGACCTT GTTATAGAGA GGCCTCGACC AGATTCAAGA CCAGCAGTTA AAAAAGAATC AACTTTGCCT CCCAGGACCT ATTGGAAAGA AGCTAGAGAG AGAGATTGGT TTCCAGATCA AGGATACAGA GGTCGAGGCC GAGGTGAATA TTACTCCAGA GGGTCGAAGC TATAGAGGTT CTTTATGGGA GGGCCGTGCC AGNGGGTTGG TAGGGGGACA CACTTCGAGA TTATCCTCAG TATANGGGC AATAAGCC

SEO ID NO:1740: (Length of Sequence = 376 Nucleotides)

GAATAAATTC GCAAACTATG CATCIGACAG AGGACIAATA CCCAGAATCT ATAAGGAACT CAAAAAATCA GGAAGAAAAA AAATCCCCATC AAAAGTGGGC TAAGGACATG ANIAGACAAT TITCAAAAGA AGATATGCAA ATGGCCAGAA AGCATATGAA AAAATACTCA ACATCCCIAA TITATIGGGGA AATGCAAATC GAAACCACAA TGCAATACCA CITTACTCCT GCAAGAATGG CCATAATTTA AAAWTCAAAA AATAATAGAT GITGGCGIGG GATGTGTTGA AAAGGGAACC ACTITTACAC TGCTAGTGGG GATGTVAAAC TACTTCGGCT ACTATAGAAA ANCAGGATGG GNGGATTCCT TAAAAG

SEO ID NO:1741: (Length of Sequence = 322 Nucleotides)

CARATGCARA RATCARGACT IGICATARAN IGIATCICCA TAGCCIATAC IGITTARATU ACINTAR IN INTESTRUCT CITGATGITT RATACAGCAR ATGITARACC RAGCITICAC TACAGRARIA RACAGRARIT TATAGGCGCI CATTATCCIT TIAGACARAG TIGIATTIGC TITGCIATIR TITTIGITTA GGNITIKIGC RACIATTICA CARACAGGNA CRAWRATATT

TAAATTGITA ATAGAARTIT CCAGITTICT TTAGICICIG GCTACTCCAA GTACTGGITG CTGTGAATGA CCTTTTCATG

SEO ID NO:1742: (Length of Sequence = 322 Nucleotides)

CCCCCCAGCC AGGAAAAAAA AAAAAAGCTT TGAGGAATGA GAGAGGTGA GATGGTCGA AGAAGGTGCT GGGCAGCCAC
GGGCGCCACG CCTCANTGGC CCCAATTGCC GAAGCCGATC TCCTGCTTGT ATCTGTTAGT GAGGATGTTG GCTTTGCGCG
TNAGTTTCGA GAGCACAGTG TGCAGCCCGC GCAGGTGGTA GTGGAACTNC TGTTCCAGGT CTTCTTCGCC GGCGTCCGAA
CCCTCCAAGT GGCCCAGGTC CACCAGGATG TCCTTGGGAC TTCCAGGCAC TGCCCTNCTC GNTCCCAAGC CGGTNGGAGG
CG

SEO ID NO:1743: (Length of Sequence = 250 Nucleotides)

ATGGTIAGGG GCCCAACGCA GTCACCGCCG TCCGCAGTCA CAGTCCAGCC ACTGACCGCA GCAGCGCCCT TGCGTAGAGC
CGCTTGCAGC GAGAACACTG AATTGCCAAC GAGCAGGAGA GTCTCAAGGC GCAAGAGGAG GCCAGGGCTC GACCCACAGA
GCACCCTNAG CCATCGCGAG TTTCCGGGCG CCAAAGCCAG GAGAAGCCGG CCATCCCGCA GGNCCGNGTC TTTCAGCGAG
ACCHGAGTTT

SEQ ID NO:1744: (Length of Sequence = 247 Nucleotides)

GATGATIGAG TGTTTCTTTA AAAATAAAAA CCCCACAAAA AAGCCAGAAC ACCCTACCCA ACCCAGCCCA GTGTAACAGG
TTAGCCATTA ACACAGNATA AAGAWGGTCC CAGCCACACA CGTCATTACT CGGCAGAGGG TGTCCAGKCT GGTCGKCCGA
CGTCACAGTG GATGGCCCTG CGTCGCTGGG RCACAGACAG GGNGCAGGCA TGGCACCTTT CGNCACGCAG AGCAAGCATA
GGCTGTA

SEO ID NO:1745: (Length of Sequence = 379 Nucleotides)

TTCTAAACCA GITAATAAAT TCATTCCACA AGTATTTACT GATTACCTGC TTGTGCCAGG GACTATTCTC AGGCTGAAGA
AGGTGGGAGG GGAGGGCGGA ACCTGAGGAG CCACCTGAGC CAGCTTTATA TTTCAACCAT GGCTGGCCCA TCTGAGAGCA
TCTCCCCCACT CTCGCCAACC TATCGGGGCA TAGCCCCAGG ATGCCCCCAG GCGGCCCAGG TTAGATGCGT CCCTTTGGCT
TGTCAGTGAT GACATACACC TTAGCTGCTT AGCTGGTGCT NNGCCTGAGG GCAGGGCAGG AAAATCAGAA TAGCATTTGC
TTTCTCTGGG GCAAAAATGG GAAAGTTCAG CGGGNNGCAG CAGGAATCAA GTGGGCATT

SEQ ID NO:1746: (Length of Sequence = 472 Nucleotides)

TTCATGCTGT CCCTTCATTG AATTITAGAA TGATTGAAGA TAGTGGGAAA AGAGGAAATA CCATGGCAGA AAGAAGACAG CTGTTTGCAG AGATGAGGGC TCAAGATCTG GNICGCATCC GACTCTCCAC CTACAGAACA GCATGCAAGC TTAGGTTTGT TCAGAAGAAA TGCAATTTGC ACCTGGTGGA CATATGGAAT GTCATAGAAG CATTGCGGGA AAATGCTCTG AACAACCTGG ACCCAAAACAC TGAACTCAAC GTGTCCCGCT TAGAGGCTGT GCTCTCCACT ATTTTTTACC CAGCTCAACA AACGGGNTGN CAACCACTTC ACCAAAATCC ATGTGGAGGCA GTCCATCAGN CTNCTNCTTA ACTTNCTGCT TGCAGCGTTT TGATNCCGGA AGGCCCATGGT AAAATTTTCA GTATTTGCTT GTCAAAAANG GGTTTTAGGC NCCATTTGTG TGGGAGGGGA AG

SEQ ID NO:1747: (Length of Sequence = 351 Nucleotides)

ACCATCAGAA TACTITAATA AGATACCAGT GTCAAAATAC ATTICCTTAT AAAGITAAGC TOCCATACAG TTATAATGIT
GTCAGTAGGA ATTCGACAAT ATAATAACGT TCATCAAATC GTTACGTTGA CAGGTAGGGT TAATAGGAAG GTTAGGAATAT
TTTCCAGTGT TITAGTAAAA CTGCAAGGGT AAAATGCCCT TAATGCCAGG GCAACACAC CAGGNAATCA AATACCAGCA

TTTACACENC AGIAACCCTT CAAGTTCTGC CACCCTGTGT GGGGGTAATG CCGTGCAGCT AAAAATATGG GTTTTACGNA ACANCCATGG CCTAAGGGGA TTTCTCATAG G

SEO ID NO:1748: (Length of Sequence = 428 Nucleotides)

AATAGCTICA GCIGATIGGG IGAGICCIAT ICATGITATA AAAGGTACTC IGCITICCIT AACAITCCAT AAATGTTAAT CACAITCIGCA AATACCTICA CAGCAACATC IAGACIAGIG ITTIGACCCAA CAACIGGGCA CAATAGITTA GCCAGCTITTA CACAITAAAAC AICATCACAC TAIGCTICIC ITCIGIGGIC ITTIGITACCA CGIATCIGIT CCAIGIGITI INCITIGITAT ATATCCIATC CIGICATATC ICCCCIATGG ITTIGIGGGAA ACIATAAGCC ITCIGGGGGG IAAAACACTA IATCITIGIT CAATIGITAA IACAICGIATA AGCAITATCAT GCCIGGGGGC AITGGITAAA CCCCCCATIT AAATACAGCI NGGCAGCAGG ATTITAAGGCA ITCCGICATG GIGGCCA

SEO ID NO:1749: (Length of Sequence = 478 Nucleotides)

GGITTCACCA TGTTGGCCAG GTTGGTCTCA AACTCCTGAC CTCAGGTGAT CCACCTCAGC CTCCTAAAGT GCTGGGATTA

AAGGCGTGAG CACNCACACT CACACCTGGC CCTCAACCAT CTCTTTCACC TTCTGCTCAT GACAGTTTAC TAGAATTTTT

TTCCCTTGAG ACTGAATGTT AAGTCAAAAA CAATAAAAAA TTGCTAATCA TTACTATGAC TCCAGAGCTA CTTGCTTCTT

TAAAAAATTCC TGAANTTATA AAATATAAAG CCAAAGCAAT GAATTTCTAA TGGTGGAATT GTAGACACTG TGGGCCCCCT

GCGCCTTGTTA TTTTCAGATG GGGCAAGGGG ATATTCCTAA CCTATTTTTA AAATCATGCC AGCCTAGATA ACTATGTGAA

AAA.AATATGG GGTGCTTAGC AAAACTATTA CCTAGCACCC CTTTGGCAGT TTTACATTAA AAATCCCTTT ATTAGGTT

SEO ID NO:1750: (Length of Sequence = 439 Nucleotides)

GACATTITAT TICCAGGITG GCACGIGIAT AAGGCACAGG GGCAAATGGC TITGGGGTCC TGGAACTGGA AATGGAGACA GGIGIGICTC AGGIGICCCT GCCTCCACCA CCCCCTAAGT GCACTTGAGA CAGGACCAGT GGIGGIGGIT CCAGCCCAGG GICCTGAAGG GINCCACTGG CTCTAGGGGA GAGCCATGGG GACAGCTCCC CAGGCGGGAC CCTCTACTCT CCAGCTACCC AGGAGGGACC CINTCCTCCT AGGGGGCGAG GCCAGCTCCA AAGTGCTTNG TGGCTCCCCA GGCTTAAGGG ACCAGNCTGC CAGGGAGGGC TNGGNTCANA GAGAGAATAG TAAGATNAGA CGAGGAGAAG CACCCCACTA GCACGGCGAT TGGANAACAC TNTCGGCGGT ACTCGTCATG TGGGTAATTT GCCAANTTC

SEO ID NO:1751: (Length of Sequence = 347 Nucleotides)

CTCTATTACT TATGATTACA CCATGGCAAT ATTCCTTTT CACCAGGAGC TTTGGACCTG CGCAGGTTGT GGCATGTAAT CACCAGGAGC ATGTAGTCAT CTGTAGAAAT CACAGGCACA CTCATGTTTG CTCTGGAAGG AATCTGTTTT CCACAATGAC TCCCCCCAGC TAATGTACAC ACTGGCATTT TGCATGCCTT CCTCACACAT GGGGCACCAG CCTTGCTTCA GAACCACCCA AACTCCACAG AGGCCCTTAA ATATGGGCTA GGGACAGATT TTCTTTAAGA AAGAGTTAAG GANGCAGCTT ACAAAGGGAC AAGGCAAATT CCACAAGTCA GGCAGCA

SEO ID NO:1752: (Length of Sequence = 297 Nucleotides)

GGATATICIA GCCATACAGA TICAATGGAA CAGAGAAGAG AAAGGAGGIT CCATIGGCAC CATAGTGAGC CATICATITG
CCCAGGGAAG NNGGIGGGGG CTAAGGGGCT AGGITTGGTC CCATGGCTAC ATTAAATGCT TGGCATGACT CCAGGGCTNC
TCTAGITAGI GGCTCCAGCA CAGIATGAGI TAGGTGAGIT AGGIGTAGGA GITTGGGGAC AAGGAAAAAG GGAGGAGGGG
TCCCTAGAGG CINGGIGCCC ATTACATAGA CTCAAATTCG TCAATGCGCT GCTTTAG

SEO ID NO:1753: (Length of Sequence = 402 Nucleotides)

AAATTIAACT TCAACAAGCT GGTGATGCCC AACTACCCAT TCATCAACAT TCGGTCAAGT GGTGTGGTTC CTCAGAGTGC ACCACCAGTG CCAACAGCCT CTTCCCGGTT CCATTTCCCA CCTCTGGACA CCCATTCTCC AACCAATGAT GTGCAGCCGG GACGGTTCTC TGCTAGCTCC CTAACTGCTT CTGGCCAGGA GTCCAGTAAT GGTACTGATA GAAAGACTGA GCTTTCAGAG CTGGAGGATG GCTCAGCTGC TGACTGGCG CGGGGTGTG ATCCCGTGTC CTCCAGGAAT GCCATTGGTG GAGGAGGGAT TGGCCATCAG AAACGCAAGC CTGACATAAT GCTTCCTCTG TTTGCTAGGC CAGGGATGTA CCCTGACCCC ACAGTCCTTC GT

SEO ID NO:1754: (Length of Sequence = 397 Nucleotides)

CAGTIGGCATC TATGGCTCTA AAATGGAAAG GAGGAGTCCT GGATTCAGGC TACTGACTTA CTCTGTGAAT TTACACATAA
CTTCCTTTGA GCCACAGATT TAGCATTCTA CCAGTCACCT GATATTTCTG AGCAGCCACA ATATTTTAAA ACTATATTTA
AATCTGAATT TGGATTTAGC AGAATTTTAT TTTTTCCATT TCTATTTTCT ATGGTCACTA AATTGAAATT ACAACCATTG
TAAAAATTTGA TATCATTAAA TATGTAGGAC TTTATCCAGT TTCAAAGTAA AGATGTCTCT AATGTAATTA ATTGTATTT
TCACTGATGA GACTGAAATA CAATCAGTCT GTATTGTGTG GTGCGTATGT ATCAGTGGTA AGAGGCTATG ATTAGAC

SEO ID NO:1755: (Length of Sequence = 353 Nucleotides)

GAATTACTCT GTTGTTCACC TTTTGCTTTT TGCACTGTTT GINCTCTTAT CTGTATTTTG AGCTTAGTGC TAGGACTGAG
AGGCTGCACC ATAGGGAATG TATGGGAGAT GGTGAGGGGT GCCAGTNAGG GGTGCGTGGA GGAGAGGCCT GGGCTCCTCT
ACTGGATCTA CACTCTGTCC CAGGTTTTTA GATCCCACTG AGCCCAGCTG ACTGAAAACA AGGACAGTCA GGGTGAAACT
TCTTTTGCCA GAAGTGTGGC CTGAGTTGAA TTTCTGGGAG GATGACGCAG ATGTCTGCTG CAGAGCTGGG CTGAGAGTTC
TNCAATCTAG CTCTGACTTA GGTCAAGGGG CCT

SEO ID NO:1756: (Length of Sequence = 184 Nucleotides)

TOGGCTCGGA GCATCGAGCT GGACATGCGC ACCATTGCCA CTGCACTGGA ATATGTCTAC AAAGGGCAGC TGCAGTCTGC CCCTTCCTAG CCCCTGTTCC CTCCCCCAAC CCTATCCCTC CTACCTCACC CGCAGGGGNA AGGAGGGAGG CTGACAAGCT TTGAATAAAA CACAAGCCTC CGTT

SEO ID NO:1757: (Length of Sequence = 425 Nucleotides)

ATTACAGGG TGANCACCAC ACCTGAGCTA ACTTCCTGGC TTTTCAATCA AACCATCTTT GTCACTTCCT GTCCCCACCT
GAAGTCAGAA AGCCTGAAGA GAAGACGGCT CTATTGCCNC AGCTGGAGTG TGGTGGCACA ATMTCAGCTC ACTGCAACCT
CTGCCTCCTG GGTTCAGGCG ATTCTCCTGT CTCAGTCTCC TGAGTAGCTG GGATTACAGG TATGCACCAC CACGCCCTGC
TACTTTTTCG TATTTTTAGT AGTAGAGATG GGGTTTCACC ATGTTGGCCA CGCTGGTCTC TATCTCCTGA CCTCGTGATC
CACCTGCCTC AGCCTCCCAA AGCGCTAGGA TTACAGGCGT GTAAGCCACC ATGCCCCGCC AATTTTGCCA GTTTTTATTG
GGCTATTCCT TATTGAGATC TAGGG

SEO ID NO:1758: (Length of Sequence = 407 Nucleotides)

AGGAAGGCAT AAGCTAAGCA TCCTTCTAAC CAGTTCCCAA AGTCCCATCT GCCTCCATGT ACCAGCTGAT CGCAGAGCTG
GACTGGGGCA GGCTGGGCTT CCAGGAAATT CCTGAAGTTC TGAAACAGCT TCCCCTCTAG AGAAGCCCAC CCAATGTGTT
TTTTAGTGAC AGGAAGAAAG GAGGGAAGAG CTGATGTGGT GTGGCCTGCC CATATCATAC AACCCCACCA GGAGCAGGGC
AGTTCCCCAAG GTGGGTGCCC GTAGATCTGG GAGGCCAGGC TGGCATGATT CCTGTGAAGA ACTGTGCCTG TNTCGTCAGG
GAGAGGCCTG AGCCCTCTCA GAAGCAGGGA CAGCCACAAC TGAAGAGCAC GCCAAGCTGA CGCAGCAGCAG GAGCAGCTG
GAGCAGT

SEO ID NO:1766: (Length of Sequence = 373 Nucleotides)

GTAAAATACT AAGACACTAA ATGCGTATTT TAAATTTGCC CATTAAGTTT TGGGCTGCGT AAGAAATTAG TAAAAAATAT
TTCCAAATAA CATGCAGAAG TTGTTTTTAA ACTTAAAATC TCATATTTTA GCTACACCCA CAGCGATGCT ATAGAGAGGA
GCTGGATTTC GTTGTATCTG AATGGCTCAG ATTATGTTCC TTCCAAAAAAA GTTATTTTAT GTACGATCAT TTTTTATATG
ANGCATATGA AAAATCACCC AGAATCTACC ACGTATTTAC CACATAGACA AATGTCCATC TTTAGATCTG TCATTCAACA
CCATGTTATT CTTTTTATGC AACAGAATGC AGTGTTGTGA GAAGTACATC AAG

SEO ID NO:1767: (Length of Sequence = 330 Nucleotides)

GETGACAGTG GCGGCANAGC AGCAGCATGG TGGCAGCCAC CAGTGGGCCT GGGGCCCCCG GGGGAGAGAGA TGCCCCAGAG GTGCATGAGC AGACCTCGTA ACCGTCCTCC GAGCGGCTCT GGTCATGTTG TCCTGGAGGG GCGCGGGGCC CCTCTGCCGC GTCCACGCCC GCAGCCACAG ATCCATCGGC CTGTGAGTCT CCACACACCA GCCAGTCCCG GGCCGTGGAC TGTGGGTACC CGGGTGCCAC CTCCAGCTCG CCATCCAGCA CTTTCCAGTA CTCCTGGCCA CGGAAGAAGT AGGAGGCACC GTNGGACCAG CGCATGGCGT

SEO ID NO:1768: (Length of Sequence = 361 Nucleotides)

AACTGGAAAA CCAAGACTGG TAGACTCTCT TTTTCTTCAG ACAATAGGCA GGAGCCAGGC GGAGTCCAGG GATTCTTGGA
ACACCTATCT TTTCTTCGGA GGACACTAAG TTCTATTTGA AGACAAAGTT CAATATGGCA ACAGGACTGA TGGGACACGA
AGGAGTCGCT ACCGTGATTT GGTGACAGTT CTTCAAAACG ACAGTMTCTC AAGGAAAGGT GGACCTAGGA ACTCCTGAAC
TTTTGGGTTG CCTTAAGTGA GAAATCAGCA TGGCTCAGGC AAGTCTCCTG GCTTGTGAAG GCCTAGCAGG TGTGAGTTTG
GTTCCCACTG CAGCCAGCAA GAAGATGATG CTGAGCCAGA T

SEO ID NO:1769: (Length of Sequence = 389 Nucleotides)

CAACTACCGC AGCGCCAACT TCAGAGAGCA CATCCAGCGC CGGCACCGGT TTTCTTATGA CACTTTTGIG GATTATGATG
TTGATGAAGA GGACATGATG AATCAGGTGT TGCAGCGCTC CATCATCGAC CAGTGAGCAG AGTCCGTGCT TGCTATCTGT
CTCATGTTAC AGAGCTTCCA TTACATATTA AACGTGAAAT CTATGACTCC TGTACCTTAC CTGTTCAACA GACCTGAAAA
TGAGCCATGG CATTGGGACA GGGTCACTTC TGACAGGGGA AGTGGGTCCC CAGGTCAGCC CTTCTCTTCC CTTTGGGCTC
TTGCCAAAGN TGTCTTCCCC TACTGTTAAN CTTGTTTGTC ACACGGTCGA GTTCGTATTG GGTTCTCGG

SEO ID NO:1770: (Length of Sequence = 394 Nucleotides)

GCAGITTAGA GGAAGCTCCT TCTGGGCAAG GTCAGGCGGT CCTCCTTCCC TCTCTCCTTC CCCTTTGTCC CAGCCTCAAC
TGACTCTGGC TGTGGGAGGT GTGGAGGGTC CTTAGGCTTC CCTCCCCAAC CTGGCCTCCA CCAACACCCC TAACAGGAGG
CCCGTGGAAG GCTCAGCCTC TCCTCCGCAT CCCTCCTCCT TCCTGCCTAT CGGAGGGAGC CAGGGTCCCC TAGGCTGACC
CTGAATCCTC TTCCTCCCTT CATGGGAGGG GGGCAGGAAT CCAGAGGAGG ATGAAGCCAG CGGGACCACA TGGCTTNGTG
GCTTNGACAA ACAAGCTCAG GGAGGAAATG AGGAGGCGNC GGCTTCAGAG ATTGCCACE CTGTTGGGCA CAGA

STATE OF THE

SEC 1D NO:1771: (Length of Sequence = 373 Nucleotides)

CAGAAAAGCC AAAGITTATT CCAGTGTTGA CAGAGAGAGG GTGAGCCTTG CACAGCAATT CTAAAAACAT GTCATCTCCT
TCACCTAAGA GGTAAGANCC GGCTGTAAGT CATGGGGTCA CTAAACCGGC CGCAGTTACA GTAAGCAGAA GAGGTCACGG
CTCAGGCCTT CTCAGACTTT CCCTGGGACA CACGGCTCTC TGGGGGGGCC CGGCGAAACC ACTCGGACCA GGAGCCATCG
TACACGGCCA CATCAGGCTT NCCGCAGAGG TAGGCAGCCA AGGNCACGTG GCAGGCGGTG ACTCCCTTGC GGCACGTGGC
AATGAGAGGC TTCGAGAGAT CCACCTTCTT GGGCTTGAAC AGAGCACGGA GCT

SEO ID NO:1772: (Length of Sequence = 281 Mucleotides)

AAAGIGCIGG GACTATAGGC GIGAGCACTI GCATCCGGCC TAGGIGGGGT TITGICCCCG TICIGCAGGA GGGAGACIGA
GGCTCGGAGG TICAGGGCCT GCTTGGCTGT ACCCAGCCCC AGIATGIGCC TIGGCCACAC TAGTCAGATC CTTCCCCTCC
CACTCCTGCC ACCCTGCTCC TGCCCTGTCC CATAATCCAG GTTGAATGGG GGTGGGGATT TINGGGAGCA ACGAGGGCTC
AAAGAGATGG AGATAGGNCT GTTGTCAGGC CAAAAGIGCA A

SEQ ID NO:1773: (Length of Sequence = 401 Nucleotides)

CTCTCTGCCA TGCTAACCAA CGTAGAAGAG AATAAGATAA AGCAATGAAA AGCAGAGTGG CACTCTGATA TATAAGATTC
TCTAAGAAAT ATAGAGTGAA TTTTGCCCAA AGGCCCTCAC TGAACTAATT CCTGAACCAA AAGAGTATTT CTTAATCCAA
AACTTTACAG TATTAGACCT ACGAATTCTG ATGATGCCTG ATCAGATGCT AGTTGTTCTC GACAATCCAT GCAGTTTTCC
AGTATGAAGG AAAGTAACAA ATATACCATG GTTATTCTTA TTTCTTTCTG AAAAATATCT AGGATATTTT ATAGTGTCAT
GTGGTAAAAAT ATTCATTTGA CANTCACAAT GAAGTATAAT CAGAAGTATT AGCAATTTTA CTTTGTTTAT CCTGTTAATC
C

SEO ID NO:1774: (Length of Sequence = 230 Nucleotides)

TCIGITAAAA AAAAGTAAAA ATGITACACA TAGGNAATAA ATGITAAAAAG CIATACITIG CCAAAATAAA GITTCAGCIG AAGGIAATGC TAGITATAAA TTAAATACAA TICIATTAAG NNCTIGCAAA AGICAAAGGA AGACGNAAA CICCCCCITIT TGGCAATTCA AAGGCAAAGA CCIGITCATT TATICITAAT TTINCITIAT ACAATCATTA TCCCCCACAG

SEQ ID NO:1775: (Length of Sequence = 359 Nucleotides)

ATTCAGGACA TAGGCATGGG CAAGGACTTC ATGACTAAAA CACCAAAAGC AATGGCAACA AAAGCCGAAA TTGACAAATG GGATCTAATT CAACTAAAGA GCTTCTGCAC ATTAAAAGAA ATTACCATCA GAGTGAACAG GCAAACTACA GAAAATCTAC CCATCTGACA AAGGGCTAAT ATCCAGAATG CTACCTAATT TTTAAAGACT TTTTCCGGCA TCTTGAAAAA AACCACCATT ATTTGACATA GGTAAAACTG AAAAAACAAA CTATTCATAA TTACAATTTG TGACACATTA TGTAGTAGCT AGGTTCATCA CATAAATTAC ATGNTACCCC AGTTCAAGTT AAATTTCAG

SEQ ID NO:1776: (Length of Sequence = 375 Nucleotides)

GGCAGAGGCT GCAGTGAGTC CAGATGGTNC CACTNCACTC CAGCCTGAGT GACAAAGTGA CACTCCATCT CAAAACCCCA
ACTCCCCCCA AAATTITTAA TITGGTTTGC ATTTCTTTGA TTATGTTTGN GGTCGATTGA GACTTGAGGC TGGCACTGGA
GCAGGCGTTC CCACCTGTCC CGTGAGGCAA AGGTCGTGGG GAGTGACCAA GTGCATCAGG GGGTGCAGAT GCCCTATTCT
GGCTCTTTCA CGCTCAGCCA TCTTAGCATA NGTGAATATA CCATGAGCTG TTTCTCAGCT TGTTTTATTT TCCTGGRGAG
ATAGATGTCA CTGGAATGGN CTTTNTCCAA GTGAAAGGCC ATCTTGTGCT ATGAC

SEQ ID NO:1777: (Length of Sequence = 327 Nucleutides)

GATAAGGGAG GAAAGGCAGG AGGAGATGAG GCCAGCCCCA CTGATGACAC CTTGGGCCAG GCCTCACAGC TGCAGGCATC

AGCCGGAAAC TCCAGGCTGC TCATGGTCAC TGGCGGTGCT GAACTGTCTC TCCACTTINT TTTGGTCCTT GATCTTGAGT

CCAATGTCCA CTCTCTTCTC AAAGAAGTTC ACCAGCACGG ACTCCGTCAG GATGGAGGCC AGGTTCTGCT CAAAGGAGAT GCACCAGTCG GTGTCGACGG TGCCGCCGAT GCGCGCTGTG GGGCTGCAGG CCTNCCCGGC CACCAGCACC GTGTCGTCTG CAAGGTCTTC ACATTGCAGG GAGCCCGTNC TGACCACCGA GTAGGAGGAC ATGGACATGT CGTCTTT

SEO ID NO:1778: (Length of Sequence = 297 Nucleotides)

CCCCCCAACT AGAAGAATAC AATTAAAAAA AGAGGCAGTA CACATGGTTA ATAAACAGAT GAAAAAATTA AAATTCACTT
GTACTATAAG ACAGGCAGAT TAAATTATTT TTACCTATCA AATTAACCAG AACAAAGGCA TGCACTITAG TGAGGATGAG
GAACATACAG ATTCACTGGT GAAAGTAAAT GTACACACAA CCTTTCAAGT TGATAGTTTG GCAGAAGTTG CTAAAAACAT
TTAAAGCTTT CATACTTTTG ATAAGGCTTT TTATTTTAGA AAACATATAA ATAAAAA

SEO ID NO:1779: (Length of Sequence = 353 Nucleotides)

CAGAAGTAAA AGATTTIWAT TGTYCTATAG ACACTTCTGA AAAGAGATCT AATTGAGAAA ATATACAAAG CATTTAAGAG
TTTCATCCCC AGAGACTGAC TGAAGGCGTT ACAGCCCTCC TCTCCAAGGC TCAGGGCTGA GAACGGTTAG CATATCGAWT
GATCAGTAAA AACATGCAAA AGTGAGAAGG AAAGGGAAAA AGGTGCATTC CCCTAAGCTG AGGGGGATGG AATTTCAGAA
CAGAGGAGGC AGGGTGGACA AGTACCAGGT GGCTCTCCCT TTCCCTCTGT GTTATCTTTC AAAACAGTTC CCAAGCTTNG
NGGAAAGCAA TGAGCTCCAC CCTAYTCAGC AGA

SEO ID NO:1780: (Length of Sequence = 428 Nucleotides)

CGGCTTCCCC GGAGCAGCCG ACAGGGCCAC AGGAGAATGG TATGCTGCTC GGCATGGAGT GAAGACCACC CCGTGTGCAA
TCTGTTCACC TGTGGGTTTG ACCGGCAAGC CATTGGTTGG AACATCAACA TCCCTGCATT GCTACAAGAA AAATAAGGAC
ACCGGCAGCC CTTAGTTTCA CTGTTTGCCA GCACAGACCT TTGATGGGTG CAGGCTTTTC TGCGTATTAA TCAGCCATTT
TTGTGAGAGT TTGACCCTGG AAAGGGTGCT TTGTATATGT TCTTTTCACA TAGTGCCCAG CTTGCATGAA ATGTACAGAG
AAATGTGTGG TCGTATTTTT TACTTTTGTC TTGTATATGT ATGGATAATT NGGGTCCCTT GGGCAGTAGA GGCAAAGCTC
ACCTCCCATG TAGCACATGA AAATGCTT

SEO ID NO:1781: (Length of Sequence = 459 Nucleotides)

ACCICAGATT GIGAAGGGCT CIGTAGGCTA IGITAAGGAC ACTAGAAATC TATIGAAAGG TITTAAGCAG AGAATTGACT
TGCTCATATT TITINCITCAA AAAGCTCAAT AGCIACAAAA CGGTCAATAG ATGGTAGCTT IGIGGGGCTG GGGTGAATGC
AATGATATTG CAAAACAAGA TATAGGGAGA CAAGAACTIT TAATAACCTA AACCAGTGGT ICICAAACTT ICCATGCATC
AGAATCACCT GGATGACTIG CGAAAACACA AATAATCAGA CITAATCCCT ACATTITCTG ATTTAGCAGG TATAGAATGA
GGTTTAAGAA TITCTAACAA GITCCCAGAT GGCGTAAGGT GICTCTCAGG GTTTTTACTT GAGCAACTGG GTGGATCCNG
IGGATCITAT GICCCINCGA GTAAGGGGTC AGGTACAGCA TICTCCCGGTC AGATTGTTT

SEQ ID NO:1781: (Length of Sequence = 420 Nucleotides)

GAAAGCACAG GAGCCTGCTT CCAAAGAGGG ACTGTCCCGT AATTNAGAGA TGCTCCAAGG CTGACCATCC TCCTTCTCCT
GCTGCACACC CAGCAGCCAT CTATGGCTGG ATTTGGAGAA TTTCTGGTCA AACCGGTGAG TATGAGGAGA GCAGGGCAGT
TGGGGAGAGA GGTCCCAGCC CAATTCTGCC CAGAGAAGCT CCCAAAAGAG AGGGAAGTGT CCTGATGAAG AGCCCATGAA
AGGGGTGAGA CCCAGGAGGC TGTGGAGATT GCTGCGGGCT CCTCTGGTCA GTAAGGAACC CTGACAAGAT CCCTAGGATG
GGGGTCCCTT AGTCTCACTG AAGTTCTTGT AACTTNGGAL TGGGGCCAGC TCANCCTCCT CTGATACCCG AGCTACAMAT
CTGGCTTCCC AMTTCTAGAG

SEO ID NO:1783: (Length of Sequence = 427 Nucleotides)

wight of

AGAGCTTAGC ATGCTGTTG TTCATGTTTT TATGTGTTTA TTTCACATTG ACTITTGCCG TGAGCTTTGA GGGAGACAAC
ACCATCACAT ATGTGTAAAT TGTAAAAGAA TTGGGAGAGA ATAGCTTTGG GAGATCATTT TCTTACTGGC CATGATGAAG
AAAGCTGTAT CGTAGGAAAA TTACTAGGTA ATTTTACTCA CTTGATAAAG TTAATTTGCA AGGTATCATT CGATTGGTAG
AGTTACCAAA ATGAGAGTTA AAGAAACAGA AATATGGTTT CAGTTTATGG TGCATTCTTA TCTTTTTCAC TGAGTCTATT
TCTGTCTGGT TGCTTCACTT AGTACTCCAA CCAGACAAGA GGAAGACAAC TATGCTAGTG TTTTAGGAAA TGGGACAGAA
TGGGGTGATT TAAGTAGGAG CCNGGGT

SEO ID NO:1784: (Length of Sequence = 428 Nucleotides)

ATGGGATACT AATGCAAGCA TTCAGTGAAA AAAGTAGATT ACCAAACTAT ACGATCCTCA TTTATTTTAA AAAGTGATAT CACCCAGAAA AAAATAAGAA AGATAAAAGA TGTTGGTAAA ATAACTAAAG AATAAAAATA TAGGGGAAAA GGTAGCCAAG GGATAGATAT TGATATCAT TTTCTTTTTA CAACTTTATT AAGTTGTAAT TTGTGTGCAA CAGATTGCAT ATATTTGANG TATATAAACTT GACTAATTTT GACAAATATA TACACCCATG AAACTACCAG TTATAATTTT AAACATTTTC ATGGCCCTCC AAAGTTTCCT TGTGTCCTTT TGCAATACAC GCAAACACAC ACACCCCACA CACAGTATGT AGGGCAACCA TTGATCTGCC TTCTGTTACA ATAGGGTAGG TTTGCATC

SEO ID NO:1785: (Length of Sequence = 414 Nucleotides)

GIAAACAGAT TACATTIGAA CACCTAAATA AGIATTIGIT TCATAATCAT TACATGCITG TITATGATIT ACAAAGATIT GGIAGAGAA AGIACAGTCC TTAAGGCATA TATATGCCAA TGCATTAAAC TACTCAGCTT TIGIGCCAGC TCAGGIGITC ATAGGAACAG GAATGIGGAA TACCAGCTTT TTACTTTAAT TATACTTTTA TGCTGAATTT TTCTTCCAGT TAAACCTTTA ATTACACTAG TATGIAAAGT AGITACTGAG AAAAATAAGT TTTTGATTTC CCTTCTGTTG GATCTGTAAC ATTTTTAAAT GGAGCTATTT AACACATGAC ATGCTAATGT TACTTAATGG GTCTCTGCAT TTTAATTTTA NGAAACACAA ACCTGGGTCA CAAAACATCT TCAG

SEO ID NO:1786: (Length of Sequence = 397 Nucleotides)

GITATICCAA CCAAAATITC CIAAGATGA AATGCAGAAA CITACAGAAT TGAGTAAAAA GACAAAAACG TAAATACTAA
ATATIGAAAA GATGCAAGIN CICCCCAAAT ACACTCATAG ATTTAATAAA ATTCAAATIT AAAGGCAATT AATTAGGGAT
GAGGCAAGAA TCIGGGAAGA AAATTAATCT GAAGTTTGIC TGGAAAAAATC AATGGGTGAA ACGAAAAATAT TTTAGGATAA
GATTAATGAG AAGTAAAATT ATTTCAATTA TAAANGTAAA ATGATAAAAT AGTTAGACCT ATATGGTACT GATGCCAGGN
ATGITATACA AAGCTACGIC AAGGCTTGAG GATAATTTIN TTGAAGATAT TCGTGGGTAT CICATTGGCT ATAAAAG

SEO ID NO:1787: (Length of Sequence = 408 Mucleotides)

TCCCACAATT GACAATATAT ATGCATGIGT TIAAACCAAA TCCAGAAAGC TIAAACAATA GAGCIGCATA ATAGTATITA
TTAAAGAATC ACAACTGIAA ACATGAGAAT AACTTAAGCA TTCTAGTITA GITTITIGTA ATTGCAAATT ATATTITINC
TGCTGATATA TIAGAATAAT TITTAAATGT CATCTIGAAA TAGAAATATG TATTITAAGC ACTCACGCAA AGGTAAATGC
ACACGTITTA AATGIGIGIG TIGCTAATIT TITTCCATAAG ANTIGTAAAC ATTGAACTGA ACAAATTACC TATAATGGAT
TIGGGGTTAAT GACTTATGAG CAAAGCIGGT TIGGCCAGAC AGTATACCCA ANCITITATA TAATATCCAG ANGGCTATCA
CACTTGIG

SEQ ID NO:1788: (Length of Sequence = 391 Nucleotides)

 TTCAGTGCAG AAGTCAGTCC AGGTGGGTTC AGGCCCATGC CACCTTCTCT GGCCTGCACA GTCCCACCCC AGGCAAGGGG TTCTTTCCAG AAAGGCTAAA TGCTCTGTCC TAANCCTNGG AAGTGTCCCT TTCAACTAAA CCCCTGGCCT T

SEO ID NO:1789: (Length of Sequence = 312 Nucleotides)

CAGGGIGAAG TGAGCCIGIG TGGGAAATGA GICTAGIGIC AGGAGGCCIG GCIGCTATAA TGATATITAT CICACAGIIT ATATITCATT CATITATATI ATITITITIAA AAGGITTCIT TATCAGCIAC TAAACATCIC AGCAATITGG TGIGCATAGC TCTAGATTAA GCAACAAAGA ATIGTACIGA TAACAAACCA CAGGGGAAAT GGIGGITAGT AAGAGICAGC CITATAAAAT TTACATCCAC ACIGITTCAC AGCAAGATIG CICICTCCAA AACGIAGCCA TCAAAAGCAG CAAACAAACC CT

SEO ID NO:1790: (Length of Sequence = 281 Nucleotides)

TGTTTCCYTC ATTAGCTGTA GACTATCCCC TCTCCTCCCA CCACAWTGTT TCTWTGATGA KTTACAAACA GAAAGGAAAT CACATTTTCA TACTAAAAAC AAAATGWTCA GAGCCTTGAT TTYTCCACTA GAWACTACAC GTACAGTTAA GAGTCCACAT GCAACACCTT AAWTCACAGA CTGAGACCTC ACATTYTGAC CTGGAGWTTC CTCCCCTTCC CCAGCCTTGG GCTAGCTTTG GCCTAGGCTC AKGTAATACT GACACCCACA GGCGCTGCTC T

SEO ID NO:1791: (Length of Sequence = 261 Nucleotides)

AGGCAAAGCA GAAAGGTGTG TTTGCCAGAC CAGCATGGGC AGCTCAGAGG GAGCAAAGCA TCCACCAGAA GAGGCTCTCC
ATTTTTTTGT AGGGCCTGAC AGTTGAGATT TGAGGCTGAG TTAACAWTGG GACCACTGAA CTTTTTTCCA ATGGAAAAYT
CACGGCCCAG TCCCACAGGA ACTTTGCGGC ATACCAAACA ACAWTGAGGA AGGAAGGGCC GGGTGGCTCT ACCAAACAKT
TCAGGTCCAC TGGGTGAWTG A

SEO ID NO:1792: (Length of Sequence = 324 Nucleotides)

CICCATCITI ATCGCTGIA TAAACATCIC TGGICTGIAC ATACATTICA TACATCGIAG GGIGGGAAGC GAGGGCCAAA GGAGGCCCA GCAGCACAAC AGCICACCG CITTCCCTAC AGCCCIACCC GCICTGIGCA AACCAAGGCC AACAGCTCCT GCIGCCTCTT CCTCCCTGGA AAAGICACTG TTATGGGGAG GGGGCCAGGG GITGAAGGAT TAGAAGGAGA TAGAGGGCTT GGTGGGGAGG CCACATNIAA GTCCTAGATT CAAACACTGA AGCGAAACAG GCAACTGGCA CAAGCAGCAA GCTTAGGCAT GGGC

SEQ ID NO:1793: (Length of Sequence = 386 Nucleotides)

ACTOTTEGGE ACCCAAAGAT GTCAGGTCCC CATACTCTGA GGAATCAGGA CACAGCCCAG TGCCTGACAC CACAGAGTGA
GGCAGCCCTT CGGGTGAGGG CCTGGGCCTC GAGGGATGGC AGCCACCACT GCCTAGGCAA ACGCACCTGG GGCTGAACTT
GGCGCCCCGGC ACTITNAGGA CGCCAGCACC AGTGGGCACT CGGAAGTGCC AGTTCTGGCC CAAATTTGGT GACCTGGGTC
AGAAGGACCT TTCAGAATGA MITGTTCCCG TCAGCAGATA CCGTCAAGAC ACGGCTGGCT CTGAGAGGGG CTGGGTGCCC
GTTTTGCCTG TATTCTCCTG GGGGCCAGCA CGTCTCAGAG GGTGTCCCTG TGGGTCCCCG GGGTCA

SEQ ID NO:1794: (Length of Sequence = 308 Nucleotides)

GGATGCTCTT TAAAACATGC AAATTGGGCC GGGCACAGTG GCTCATGCCT GTAATCCCAG CACTTTGGGA GGCCGAAGTG
GGTGGGTCAC CTGAGGTCAG GAGTTCAAGA CCAGCCTGGC CAGCATGGTG AAACCTCATC TCTGCTGAAA ATÁCAAAAAT
TGGCCAGGCG TGGTGGCATG TGCCTGTAAT TCCAGCTACT CGGGAGGTTG AGGCGGAGA GTTGTTTGAA CCCGGGAGGT
GGAGGTTGCA GTGAGCCGAG ATTGCACCAT TGCACTCCAG CCTGGGGTGA CAGAGCGAGA CTCTGTCT

SEO ID NO:1795: (Length of Sequence = 418 Nucleotides)

GAAACGCTAA GGITTIGACA GCGITACAGI GAATICICCG GCTGIAGAGA TIGGAGGAAG TCGGGGGAAA TICGICICTA
AGITGIAAGG TGGAACAGCA TICATITICI TACIGCCAAI GGAGGITITI CATGAATITA CIAACTCAGI AAAAAGATIC
GGCTTTTITI TITIAATCIT AAAGGATCAC GCITTAAACC TCIGIAACAA AGIAATTATI TGIACCACIC TCTACCCCAC
CCTCCAACAA AATAACCIAI CGGNICICAG AAAATAATAA CCCTTIGCCT GCCTITGAAA TAGITATCCT TITIAGIATG
ACAGIGITCA AAAATTCITI TCTTAGACIT GIAGCAGAAC ATAGCTATGA TGATCTGAAT TTTTCTCTTT CAGCTTGTTC
TAAGACGAGG GGGACTCC

SEQ ID NO:1796: (Length of Sequence = 416 Nucleotides)

CTITATIACA TATECAACCT TECCATECCT GCCAGITAAC TCCCCTCCCG CCAATGITAT CCTCATGATA TCAGCTCCCT
CTTGGGGCCA CTGAGCTGCC CCCCTTTCCT TCTGGGCTGG AGTAGTGGTG CCCCTCAAGC AGGCAATGGG CAGGGGGAGA
TCCACAATTA ATCGTCGCAG TTCTCTTAAA AGTATTAACA CTTAAATAAG CACTCTTGGG GAGTTGCAAA GGATATTCAG
GATGGGATGC AGTGGGAGGC TACCCCTCAT CCAAGGTACA GGCTGGAATG AGCTACAGCT GGGTCTATCG TGGGCCTCAG
AAGGTGAAGA GGGACCCTAT TCTGGGGCTT AGTGTGGGTG GGGCATATCC TCCCCAAACT TGTTCTGTGG GCGATGTTCT
TCACATCTAG GAGAGC

SEO ID NO:1797: (Length of Sequence = 298 Nucleotides)

AGGAGGGAAA CCAGAATCAA ACTACTACTT CTAGATGAAC ACAGGCTCTT GAGAGTCCCC AAGAGAGGAG GCTGTTGATC
CAATCCTGAC TCAGACTACC TACCTGGCTT CCTGGCCCTA GGAGGTAATA ATGATAGTNT CAGGGGGTCC ATGTAGCAAT
CCAAGCAATT CCTGAGGTGA GAGCAAGCAA AGAGGATAGG ATGAAGGGAA GGCAGGCAAA GAATGTGCTC CTAGTAAGAA
GCAACTCINT TCCACTCACT TCCTTTTGCT CTNTGGCAGG CAAGTCAACT GGGTTCTC

SEO ID NO:1798: (Length of Sequence = 245 Nucleotides)

CTGGTCCATT TTTACAACAN ATACATCCAA AACACTATAT AATANNITIT TTTACAACAT TTCCAAATGA GAAGATTGCT
TTTNCCCCCA CTACTGCTAT TCACACACAG TACTTCCACG GCACAATACA TTAGGAGATC TAAAANTGCT CACCCTGTAC
TCTAGGCTGC TTAGGAAATG TGAAAACTAG NAACATTTAT AATGGCATTA GCTCCTTTCA ATACAAGGCA ACATTTTAGN
AACCT

SEO ID NO:1799: (Length of Sequence = 312 Nucleotides)

GAAATGTTAG GCTAGTTAGA AGGACACGGC AATAGCCTTG AGATTYTCAA CCAGGGTAGT GTATTAGAWG TAAAAAGGAG AGGAAAGATT TGAGAGTTAT CTCAGAAACA GAACCATCTA ATTTTTTTGG ACTGATTTGA CTGCTCTTTC ACTCATTTTT TTATTCACTC AACAACTATT TTTGAKTGNT TTGGATGGGT CAGACATTGC GCTAAGTGAA AAATAGGAAG GTAAGAAAAA GAAGACTCTG AAGATGAATT CCCTCCCCCAA AACTGAGCTA CTAGCTATTA CTCAGTGGGG CTGAAGTGAC AC

SEO ID NO:1800: (Length of Sequence = 309 Nucleotides)

GECATGICAC ACTAGECCAC AAGCGATAAG CACAGGCACC TGACTITIAA GITTITIGITT GITTGITGIT TCCCAAAGTG CTGATAACAA TAACAACAAC AATAGGATTC CAACCAGGNG CCTCAAGTGA CAGCCAGGNA GAGACCTGAA GGITGGGGCC ACCACAATGC CAAATCGITT CTAAAGGAAG CTGAAAAATG GGACTGTCTT TTGCCCACTT CGITGIGTTA AAAGGGGACA TTTGINCAAA CTNCCCAACC GAGTTCTAGA AGNTCCTGAC AAGGAGGCAG CATCCAGCCT TGACCAGGC



side for

SEO ID NO:1801: (Length of Sequence = 166 Nucleotides)

CAAAANITAC TCTGCAAAAT TAATATATGA TTTACCTGCT GTTNTCATAA GATTTCCAAA TAGACAAACT CGGTATGCTT NGGATTTGCT TTACATTCTA AGTGGATTTG GAGGTTCAGG CAGGCGCCAA GGAGTNAGCC GAAGTTTCAT CANGCGGAGA TGTTGG

SEO ID NO:1802: (Length of Sequence = 281 Nucleotides)

GGIGGATGIC TITIGGGCGCA GGATGGAGCC CAGACCCAGT GGITACAGIG TGGAGCTCTC TCCCTGTCCC CTGACTCTGG
CCAAGGAAGT GAATGCAAAG CAGCAGGGAG GAGGCAGGGT GGGGACGGCC CTCTGAGCTC TCCCGCGATGG CTGGCGTGAG
GTGCCTCTAA GACTTCTNGG CAGCCCTGCC TTCCCTACTC AGTCTTCCCG ATCTTNTTGC CACCTTTCTG TGTGGGCCAG
NCTCCCGCCA GGTACTCAGA GGCCGCTCAG AGGGCAGGGT T

SEO ID NO:1803: (Length of Sequence = 429 Nucleotides)

TTCACAGITA TAGITGEGGA CATTAACAAC CCTTTCTCAA TAATTGATAG ACTACTAAAT AAAAAACCAT GAAGGATATA
CAAGAACTGT ACAACACTGG CCGGGTGGG TENCTCATGC CTGTAATCCC AGCACTTTGG GAGGCTGAGG CGGGTGGNTC
ACTTGAGGTC AGGAGTTCGA GACCAGCCTA GCCAACATGG CGAAACCACA TCTCTACTAA AAATACAAAA AAATTAGGCT
GGCTGGGTT GGCTTAATGC CTGTAATCCC AGCACTTTGG GAGGCCAAGG TGGGCATATC ACCTGAGGTC AGGAGTTTGA
GACCAGCCTG AAAAACATGG TGGAAACCCA TCTCTACTAA AAATACAAAA ATTAGCTGGG TGINGTGCGT CTGAAAAAAAT
TAGGTAAACT CCGTCTCAAA AAATAATAA

SEO ID NO:1804: (Length of Sequence = 278 Nucleotides)

GACCTEAAGC TCAAAGTCTC TCTCCTTACA CAACCAGCEN CAACAGGGCC AAGCTACTGG CTAAGAACAG ACAAACTTTC
CTGCTTCAGA CCACAAAGCT GACCCGTNTT GCCAGACGCA TGTGCAGGGN CCTNTTACAG CCAAGGAGGG CCGCCCGACG
GNCTTATGCT CCTATCAATG CCAATGNCAT CAAAGCAGAG TGCTCCATTC GNCTTCCTAA GGNCGNCAAG ACTCCATTNA
AGATTCACCC TCCTGGTGCG GCTGNCCCTG GGAACTAT

SEO ID NO:1805: (Length of Sequence = 349 Nucleotides)

GCATCCATGG CGGAGGGCGG CAGCACGACG GGCGGGCAGG GCCGGGCTCC GCAGGTCGTA ATCTGAAGGA GTGGCTGAGG
GAGCAATTTT NIGATCATCC GCTGGAGCAC TGTGAGGACA CGAGGCTCCA TGATGCAGCT TACGTCGGGG ACCTCCAGAC
CCTCAGGAGC CTATTGCAAG AGGAGAGCTA CCGGAGCCGC ATCAACGAGA AGTCTGTCTG GTGCTGTGGC TGGCTCCCCT
GCACACCGTT NCGAATCGCG GCCACTGCAG GCCATGGGAG CTGTNTGGAC TTCCTCATCC GGAAGGGGGC CGAGGTGGAT
CTNGTGGACG TAAAAGGACA GACGGCCCT

SEQ ID NO:1806: (Length of Sequence = 403 Nucleotides)

GTGCAGTGTG GCCAGATCTT TTCTAGTAAA ATGTGTGTTA CTGATGGGCA GACAGCTCTC ATTCAAGCAG TGACAGATGT

AAGCNCTTCC CATTTTTGTG GCCCCATTGT ATTCAGCGTG TGGCTTCCAA GTTGCCTGGG ATCATCTCCA CCCAGACTAA

GGAAGAGGAA AGAGCTTGGA CAACTGCACT TGGCTGGTTT TNATGGATCA GGCAAGGAAT TGGCTCCAAC ACATTAGCTC

ACATTCCATT GGTTAGAACT GGGTTTCTCA ACTATTAGTA CAGGGTGAGT GTAGGGTTTT GGCACCATGG GCATTTGAGC

TGGCCAAAGG CTAATCAGAG TTAGAACAAA GCCACAAAGC CTGTGAATGG TGTTTATTGT TGTGAGGAGC TGTCTTGTGC

ATT

SEO ID NO:1807: (Length of Sequence = 426 Nucleotides)

GTCCTCAGCT TCACTCTGGC ACCACTGTGA GCACCGGGAA ACCTACCAGA AGTTGCTGGA GGACATCGCT GTCCTGCACC
GCCTGGCTGC CCGCCTCTCC AGCCGAGCTG AGGTGGTAGG CGCCGTCCGC CAGGAAAAGC GCATGTCGAA AGCAACGGAA
GTGATGATGC AGTATGTGGA GAATCTAAAG AGGACGTATG AGAAGGACCA TGCGGAGTCA TGGAGTTTAA AAAGCTTGCA
AATCAGAATT CAAGCCGCAG CTGTGGCCCC TCTGATGGG TCCCTCGCAC GGCACGGTCC ATGTCCCTCA CGCTGGGAAA
GAATATGCCT CGCCGGAGGG TCAGCGTTGC TGTGGTTCCT AAGTTTAATG CCCTGAATCT GCCTGGGCAA ACTNCCAGCT
CATCATCCAT TCCTCCTTAC CAGCTT

SEO ID NO:1808: (Length of Sequence = 431 Nucleotides)

GGIACITITC CATTAGATT CAAATGAGC TAAAATTAAG AGITTTATGA GCTGITAAGA ATGAGGIAGI TICICCIAGG
ACCCCCAAA GACAGTGCAA GIAATGACCG TITGGMCTC ATTCGTCGAT CITTGATAGI ATGINCIGGA GICIACICCC
CAGGAGCCAG GACAGGCGIG AAGATGGAGI CCTTGTCGCA GCTGGAGCCT TGCCTAGCTG GIGATCACAC AGCCTGGNCT
GIACCCGCAC CCCACTGGAT GGTGGTACAT GGTGGCAGGG ACAGGACCAC ACCCAGTTAA GGCCAGACCA GGCTGAGTGT
GACCCCTGAG GIAAACACTT CACTAAGCTG TGTCTTGTTC ATGCCCCCTG CTCAGTGAAA GGTGAGTCCC GAGACCAGTT
GGGTACCTCT CTTATGCGAA CCAGAGACAT T

SEO ID NO:1809: (Length of Sequence = 401 Nucleotides)

OGTGAGGCCT TGAGCACAAG TGCAAGCGGG ACATCCTGCT CGGCCGGCTC CGGAGCTCGG AGGACCAGAC CTGGAAGCGG
ATCCGGCCCC GGCCCACTAA GACCAGCTTC GTGGGCTCCT ACTACCTGTG CAAAGGAGGA GATCGACGTG TGGACCGAGG
AGCGGAAGGG CACCCTCAAC CGCGACCTGC TCTTCGACCC GCTGGGGGGT GTTAAGCGCG GCAGCTCACC ATCGCCAAGC
TCCTGAAGGA GCACCAGGGC ATCTTCACCT TCCTCTGCGA GATCTGCTTT GACAGTAAAC CCCGGATCAT CAGCAAAGGC
ACCAAAGGACT CTCCGTCTGT MTGCTTCAAC CTGGGCTGCC AAGAACAGCT TMTTACAACA ACAAGTGCCT GGTGCACATC
G

SEO ID NO:1810: (Length of Sequence = 233 Nucleotides)

AAGTGCTATA TTCATTGTAT TATAGAGAAG GTTGGGGAGC ACAGAAGAG ATCAACCCAG CTTTAGAAGG ATTAGAGAAA GCTTCCAGAG GGGTGGACAT TTGAGCTAGC AAGAAAGCAC AAGGGAAAAG GCATTTAGAC AGAGGAGACA ATTTGTCCTG ACCCAGAAGC ATTGGGGTAT GCTATGCATG GATAGNCAAA GAATTTTTGC AAAAGGGGGG CCAGCAAGGC ATT

SEO ID NO:1811: (Length of Sequence = 423 Nucleotides)

SEO ID NO:1812: (Length of Sequence = 394 Nucleotides)

 AGAGACTGAT ACCACTGGAG TGGCTGGCTT TTAATTCCCC TGGGCCAGAC CTGCAGCCTT GCCTTAATCC TTAA

SEO ID NO:1813: (Length of Sequence = 344 Nucleotides)

SEO ID NO:1814: (Length of Sequence = 442 Nucleotides)

GACACAGCAG GCCCTGCCC CTGAAGGAGA CTGCATTGGA ATTTTTGCCA GGTGGCCCTG ACACATAGGA ATGCCCAACT
ACTGTGACTA CCCTCTGAGA TAAAAAGCTG TCCTACTGAT TTTAGAAGGC CAAAATTAGA GGTCATTTTG GAGGTCATGC
CAGTGGACAT ATAACAGTTT GAAATGCTTG TTCCCCGGTG CCGTAAAGAA ATAGTACTTG AACTTAAATT TATTCAGCAA
GGCCATTTTT ATTTTCTGCA GAAAGGGTAC ACTTGGCAGC AGTTTTNCCA CGAGAGTACC CCGAACAAAG GAGACAGGGT
CATTTATAAC CTGACGCGTC CACCCTTCTG CTGTGTCCGG TTTCCATTGG CTGGAACAGG ACCTCACATT CTGTATTTGT
CCCGATTGGC TAGCAACTTA GGACTTATTA AAAGAGGCAA AG

SEO ID NO:1815: (Length of Sequence = 299 Nucleotides)

GCAGAGAATC CCTTGAACCT GGNAGGCGGA GGTINCAGTG AGCCGAGATC ACGCCACTGG ACTCCAGCCT GGACAACAAG AAGGAAACTC CATCTCAAAA AAAAATTGAAA AAAAATTCAA GANATACAGA ATGCAAAANG GGACCAAAAA AGTACCAAAAA ATTTCAAAAAT TTTGTTAAAC TGTACCAAAAT CIGGNTACGA AGCGTTATTT TTGCCCACAG GGCACTTCCC TGGAAAGNCG TTACAATAGC TNAGGCTTCC TCTTCAGATA GANTTAGAGT GGCAGTAGGA TAGGCTCTT

SEO ID NO:1816: (Length of Sequence = 286 Nucleotides)

SEO ID NO:1817: (Length of Sequence = 320 Nucleotides)

GAAAGGAAGG CCAGGGIGGG AGGAAGGATC AGCTAAATCI GAGGGAAGAA GAAGGAAAGG AGAGGACTA TIGCATAGCA GATGCAAATG AAGGGACIGI CITATTATAC AGITTIATCA TCIGITAATA CTCATAATCI TGITICTITI TCAACITITA TATAATTITA TCITTACATI AGITAAATCA AAAATCITAA AACACATTIT AAACGIGGIC ATAGGITACI TITATATATT ATTGAATTITA TAATAAACAT GITICTITINC TGGAAACIGG GATGGNACCN CGATGGIGIT TCITGAATAT AAGAGIGICC

SEO ID NO:1818: (Length of Sequence = 356 Nucleotides) 9

CCCAGGAGGC TGAGGCAGGA GAATCGCCTG AACCCGGGAG GCAGAGGTTG CAGTGAGCCG GGATTGTGCC ACTGCACTCC AGCCTGGTGA CAGAGCGAGA GTTCATCCAG ACACACACAT ATATATATAA TINCCAAACA GGCTTTACTA AACCCCCTGA GGTCTCATGA CACAGTAGAA AATCATGATT TAGTAGAAAG AGCATGGTCG TAGGAATCCA GTAGATCAGT AGACCTCTGA TAGAGTCCCA AATCTGCCAC TTTCAATCTG TATGGCCTCA GGCAAGTTAC TTAANCTTTC TGTGCTCTGTG AGATTCATTAAAAATGGGGG ATAATAATAG TAACTTCTTC ATAGGG

SEO ID NO:1819: (Length of Sequence = 328 Mucleotides)

CCACTCCTGT AACCTGCTGG ATGACTCTGC ACTGCCCTTC TTCATCCTCA CCAGTGTCCT GGGTATCCTA GCTAGCAGCA
CTGTCCTCTT CATGCTTTIN AGACCTCTCT TCCGCTGGCA GCTCTGCCCT GGCTGGCCTG TCCTGGCACA GCTGGCTGTG
GGGCAGTGCCC TCTTCAGCAT TGTGGTGCCC GTTTTGGCCC CAGGGCTAGG TAGCACTCGC AGCTCTGCCC TGTGTAGCCT
GGGCTACTGT GTCTGGTATG GCTCAGCCTT TGNCCAGGCT TTGCTGCTAA GGGTGCCATG CCTCCCTGGG NCACAGACTG
GGTGCAGG

SEO ID NO:1820: (Length of Sequence = 359 Nucleotides)

CCACCATGCT CTGCACTCGC NCTGGTACCA GGCCCGCGAC CTCATGCTCA TGAGCCACTT GCAGGACAAC ATTCAGCATG
CAGACCGGCC AGTGCAGATC CTTTACAACC GCACCATGGT GCAGCTGGGC ATCTGTGCCT TCCGCCAAGG CCTGACCAAG
GACGCACACA ACGCCCTGCT GGACATCCAG TCGAGTGGCC GAGCCAAGGA GCTTTTNGGC CAGGGCCTGC TGCTGCGCAG
CTTGCAGGAG CGCAACCAGG AGCAGGAGAA GGTGGAGCGG CGCCGTCAAG TCCCCTTCCA ACTGCACATC AACCTNGAGC
TGCTTGGAGT TTGTTTTANC TGGTGTCTGC CATGTTCCT

SEO ID NO:1821: (Length of Sequence = 208 Nucleotides)

CCTGGGTCTG TGACCCAGAG TTCCAACACA AAGACACTTT GTACTGGAAC GCTGGAGCCA TTCCAACATG AACAGCAAGA ATAGAACCTG TGCTGGCTGG TCTAAGATCA AACCTCGNGA TGGTGGTTTG AAGINCTTCT TCAAAGAAAG CTTGAAAATG AAATCTCAGT TAGGCAAGNC AGATAAAAGC AGAGTTATTC TGGTGGCG

SEO ID NO:1822: (Length of Sequence = 314 Nucleotides)

GGATGINITG AGCCAGAGIT TAAGCCIGAC ACACAGGCIT TGGICCICAC TGAGCIGICI CCAAGACIGG AACTACTIAG
TGACICGGCA AATTITCIGC CCCCCACCCC TCATCAAAGC TGCTAGITCA GATGITGACA GIGITTICAT GAATGITGGA
ATCITACIAG TCCAGACITA CITAGGATGI TGITGGGGAA GGCACTIGGG AITTICIGIG TCITGCATIC ACAGAGGGAG
GCCATTICAG ATTCAAGAGC AITKGATTAG GGGATCGIGA GGCAGGGATG CTACTGCGKA TITCICICIT CAGG

SEO ID NO:1823: (Length of Sequence = 344 Nucleotides)

AACAATTITG TCTTTACTAC ATCTTAAAGA ATTAGAACTI GGGTTGGTGT AAGTGACTTA CTTCCAGGGN ATCATGCTCT
ATTTCTACCA GCAGGTCATA CCCNAATGTC ACACTATCTA TTGTTAACCA TGAATGNTAT TCAGATCTAT TACTTTTCGT
GAAAAGTGGA ACATGTTACT TCCAACCATG GCCTGTCACC GTGAGTGTGA TCANCTTTNT CCAAAACCAC ATGGGTCGCA
GGAGCTAAGG GGTGGTACCC MAATGTTAGG GACAGTGTTA GGGAAGGGCA AGGGAAAAGA AGTGACTNGA TGTCTTATGA
GRAACCCGTA AATGGCTTAA AAAA

SEO ID NO:1824: (Length of Sequence = 340 Nucleotides)

GTGAGTGGCA GGTATCATGA ACCACATTGT GGACCTGGAG TTGCTAGGAC CTTTTCTGCC ATTACACAGA AAAATCCTCC CTGAGAACAC AGCCATTNGA GGNCACATGG CAGAGGAAGA TAAGACAATA AACAGAGNCA CATAATTATG GCCAGCGTGG GGGCTNACGG CTGTAATCCC AAAACTTING GAGGCCGAGG TGGGCAGATC ACCTAAGGTC AGGAGTTCGA GGCCAMCCTG GGCAACATGG TGAAACCCGT CTCTACTAAA AATACAAAAA TTAGCCSGGC GTGGTGGCAC GGGCCTGTAG TCCTAGCTAC TCAGAGGGTT AGGCAGGAGA

5FO ID NO:1825: (Length of Sequence = 357 Nucleotides)

AATTIGGITG TEGCCAAATT CTCAGTCCAA TCACCCIGGC CCAGGGCCIG GCGTGGGAGG ATGTGGCAGG CTCTGTCTCC
TTCIGGGGIT CCTGGTCTGG AGGAGTCTCC CCAACAGCCG CAAAGCTGGC TGTTTTCCGC CCAAAGCCCC AGAACTITGA

and the second s

ATGAGAGGCA AATCTACCCT GAATGCACCT CCCTCCTAGG CTGGGTGAGG TCACGCAGAC ACAGAAGGGC AGGACAGAAC TCCCCCATCIT CTGGGGGCCA ATTCGTCTGG ACACTGTGCG GTCANCITCC TTTTTAAAGT GCCAGTATCG GTGGGGCAGG AAGGGACTCT CAGGGCTGAG CAGAGCCTTC TTCAGCG

SEO ID NO:1826: (Length of Sequence = 207 Nucleotides)

COGGCCCCTT CAGTCCCCAG CCCCTGCCCC AACTCCGACT CCTGCACCCA GCCCGGCTTC AGCCCCGATT CCGACTCCCA CCCCGGCACC AGCCCCTGCC CCAGCTGCAG CCCCAGCCGG CAGCACAGGG ACTGGGGGGC CCGGGGTAGG AAGTGGGGGG GCCGGGAGCG GGGGGGATCC GGCTCGACCT GGCCTTAGCC AGCAGCA

SEQ ID NO:1827: (Length of Sequence = 309 Nucleotides)

SEO ID NO:1828: (Length of Sequence = 382 Nucleotides)

ATCTCTGACC ACCCCTCCT CCCCATCCCA CCCTTTGGTA ACTCCCCCGC CCAGGNCACT GCCCAGATAT ATTCTTCTCC
TTGGGCAAGA AGTTCTGTGC ATGCAGGTCA AATCTGAAAG GGNCATTTCT TTCTTTAATG AGTGTCAGGG ATGGGGGGATG
TGGCTGATGA TATAAGGGGC CCTCCAATCA GACTTTCTAA TCTAACTGAA AAGNTAATTA CAATGTTGAT GCTAAAAAAG
AAGGTTCTGG CAAAATAGAA CTTCTGAAGC ATCATAAATC AGATGACTAA TATTTGTGAT CCCCNTTTAA ATTTTCATGT
GAAGAAGAAT AGGGGATGTA ACTGAAGRAA TGNACTAAAA GTTCTTCTAT GTATTGATAA CC

SEQ ID NO:1829: (Length of Sequence = 361 Nucleotides)

GEGEGEGET CTEGASCTEG ATGTCCAGGC TEGEGEGECT GCTGGGCCTC GGGCTGCTG TTGCGGGCTC GCGCCTGCCG
CGGATCAAAA GCCAGACCAT CGCCTGTTGC TNGGGACCCA CCTGGTGGG ACCNCAGCGG CTGAACTCGG GTGGCCGCTG
GGACTCAAAG GTCATGGCGA GCACGGTGGT GAAGTACCTN AGCCAGGAGG AGGCCCAGGC CGTGGACCAG GAGCTATTTA
ACGAATACCA GTTCAGCGTG GACCAACTTA TGGAACTKGC CGGGCTGAGC TTTGCTACAG CCATCGCCAA GGCATATCCC
CCCACGTCCA TGTCCAGGAG CCCCCCTACT GTCCTGGTCA T

SEQ ID NO:1830: (Length of Sequence = 180 Nucleotides)

AAGAACGITG GCTGCCTGCA GGAGGCGCTG CAGCTGGCCA CTTCCTTCCN CCANCTGCGN CTCGGGGATG TAAAGAACTG
AGTGGGGAAG GAGGAGGCTC CCACTGGATC CATCCGTCCA GCCAAGAGCT CTTCATCTGC TACAAGAACA TTTGAATCTT
GGGACCTTTA AAGAGCCCCT

SEO ID NO:1831: (Length of Sequence = 335 Nucleotides)

AGATCTICIA TATICCGACT ACTGATICAA ATGCTAATCC TGGACGGCA TGGTGGCTCA CACCTGTAAT CCCAGCACTT
TGGGAGGCTG AGGCTGGTGG MICGCCTGAG GTCCGGAGIT TGAGATCAGC CTGGCCCAACA TGGTGAAACC CTGTCTCTAC
TAAAAATACA AAAATTTGCT GGGCGTGGTG ACATGCGCCT GLAATCCCAG CTACTCGGGA GGCTGAGGCA GGACAATCAC
TTGAACCCGG GAGGCAGAGG TTGCAGTGAG TTATTGCACC ATTTCACCC AGCTGGGTG ACAACAGGAATTCCATCC
CCCCACCAAA AAGCG

403

SEO ID NO:1832: (Length of Sequence = 337 Nucleotides)

GIATTIGGAG AIGGGACCIT IGGAAAIGCI TIGATIAGGA AGAAGGAGCI TICAIGAACG GGATIAGIGC CCITAIAAAA
GAGGACGCAG AGAGCICICI CACACCITCC ACIGICIGAG GNCACAGGGA GAAGGCCCIG ICIAIGAACC AGGNAAIGAI
CCCCAACCAG AACACCIIGA ICIIGGACIN CCCAGAIGCI CCANAICINI GAGAAGCAAA ITICIGIGCI ITIAIAAGCIA
ICCAAAIGIAI GGAAITIING IACAGCAGCC CCAACAGACI AAGNIAITAA IAAAAIAAAG AIGIAAGAIC ICIGIIGAAA
AIGCACAAAI AAIAICI

SEO ID NO:1833: (Length of Sequence = 244 Nucleotides)

TCTCTCATTG TAAGCACAAA TTGTTCCGTG TCTGGTTATT AAAATCGCTT TGGGTCTATA ACAGCCACTC TTGTCCCCCC
TTTTAATAGA AAATTGTCAT TCTAGCCTGG ATTTCTCCCC ACTGGAGGTG GAGGGTGGGA AGAGAAGGGA GTCAGCTCTG
ACAGCTTACA AACTGGGAAG TTCTGTGCAT CTCCAGGGAT TCCAGAGTTG AAGATCTGGT TGTTGGAAGC TGGGCGCCCCA
GTGC

SEO ID NO:1834: (Length of Sequence = 322 Nucleotides)

TCCTGTACTA CACCTITGCC AACATGGCCA TGTTGAACCA CCTGCGCAGG CCCCCGTCCT GCAGTACCTG TACTACCTGG
CCCAGATCGG CATCGCCATG TCTCCGCTCA GCAACAACAG CCTCTTCCTC AGCTATCACC GGAATCCGCT ACCGGAGTAC
CTGTCCCGCG GCCTCATGGT CTCCCTGTCC ACTGATGATC CCTTGCAGTT CCACTTNACC AAGGAGCCGC TGATGGAGGA
GTACAGCATT GCCACCCAGG TGTGGAAGCT TCAGCTCCTG CGATATGTGT GAGCTGGCCC GCAACAGAGT GCTCATGAGC
GG

SEO ID NO:1835: (Length of Sequence = 178 Nucleotides)

ATGAAAGCAC AAAAGAAGIC TATCAAAATT ACAAAAACIT AAAACCGAGT AAACAAAACI TCAGAAAGAA TGAAAACAAT TGGAAAAATAA CITCAAGAAA AAAATGTAAA ATGGAAAACAA TACAAGANCA ATTTGTGCCC TCTGAAAAAAC AGAGGTTAAA GTCAGAATTT TTTTGTNC

SEO ID NO:1836: (Length of Sequence = 377 Nucleotides)

CGCCTGENAC CACACCCAGC TAATTTTTGT ACTGTTAGCA GAAACAGGGT TTCATCACGT TGGCCAGGCT GGTCTCGAAC
TCCTGACCTC AAGTCACCCA CCTGCCTTGG CCTCCCAAAG TGCTGGGATT ACAGGCATGA GCCACTGTGC CCGCCCTTTA
TGCTGAGTTT TAAGGGCTGT ATGAGACACC AGGTGGTGGG AGGGAGCTGT TTTGAGAGCA GGGAATTTAG GATACTTAGG
AAATTAGAAA ATTAGAGAAG TCATAGGATC TTGGAACTAA GGGAGAACCT TAGAGTCCTG TGGAGCAGAA CCCAGCATTT
GTATGTGGAG GAAACGGAGG GCCCAGAGAA GTTGTGACTT ATNCCGGGGT CAATCTT

SEQ ID NO:1837: (Length of Sequence = 388 Nucleotides)

GGAGAGAACA AACCCICTTA CIGGCCTIGG GCCCATCCCI CTITCTCCCA CACTGCTACT TITGAGITAT CICATITIGC
TCCCAATAGI CAGCCTIGAC TITTCTGGGC TTACCTGGGC ATCAGGGACC CATGITGCAC ATTCAGITGI CCCGATTATG
TCTGCCTIAG AGCGTCTCCT AGGGCAGCCA GTCTGGAACA GTCAGTCACC TAGGGTCCTG GAGCTCCTGC AGTCTGCCAC
TCGCTNCTTC TGCCTGATAA CAAATACTAT TCCTTTTATC CTTGCAACTC GACCCAGAAA GAGGTGGCTG TCAATGTCCA
AGGCCCCTGG GAAACGAAGG ACTGGAAATN TGAAACCACT GGGCACAGGG GGAATGGGTG GGTCTGAG

SEQ ID MO:1838: (sengt) of Sequence = 369 Nucleotides)

TCTCTTTATG CCAACAATTA ACTGGGAGCT AGGTTAAATT ATTTGGCTAG ATAAAACTAC CAGCTAGATG GATTTATTTG GTGCCCTCAT ACAGAATGCT GTAGAAAATG TAAAGAAGAG AAAGCTCCTT CCAGCTAGAA GCACATGGGA CTGCTTCTAG

THE RESERVE OF

GATGGAAACA AGTOCTGCTA TTTTCACAAT CCCTAAGNGT TCTCCAGGCC TCTGGAGAAC AAAGTAAAGT TGTAAGATCC CCAAAGACAC GGAAAATCCT GGACGAACAG ATTAGAAAATA ACTACAAAAA ACAAGTTTTT TACTTTCGAA AAGGGTACTG CACTGAAACC AAGTTGGACT TTTGGTCCAC CCCCAGGGCC CTCTTCAGG

SEQ ID NO:1839: (Length of Sequence = 359 Nucleotides)

CNNGTAGGGA AGAGGACTIT ATTGGGATGT TAGTAGGGAA ACATGAGAGG GTGAATTCCA GGGAATAGAC ACTAGGACCA
AGGTGGCGGT CACCTTAAAG AGCCATAAAT AAACTTAAAA AATTAAGGTG AGGAGGTGCC ACGTGGGGAG GCTGCTGGGA
CTATCTGGGA ATTCTTAGGG ATGGAATTTT GGAATTGGAA AGGGGAAATA AGAATTTCCA GCCGTNTCAC AAAAGGGTGT
GAAATGATCA CTTCAAGACT CCCTGCTGCC CTAGGCTGGG AGTTGGGGTT CTGGGGCTCC AGGAAGAGG GAGGTCTGGG
CTCGGCTTNA AGGGGTGAAG AGGGCCCGGT CAAGGTCGT

SEO ID NO:1840: (Length of Sequence = 360 Nucleotides)

CCAATGAGCC CAGCCTGACA CATATGGACT GCTCGACAGG TCCACTGTCC CACGAGCAGA AGCTGTCACA AAGCTTGGAA
ATTGCCTTGG CATCCACCCT TGGCTCTATG CCCTCCTTCA CGGCACGGCT GACCAGGGGA CAGCTCCAGC ACCTTCGCAC
AAGAGGGAGC AACACTTCCT GGAGGCCTGG CACCGGCTCG GAGCAGCCTG GGAGCATCCT GGGCCCCCGAA TGTGCCTCCT
GCAAAANAGT ATTTINICCC TACTTCAAAA AGGAGCCGGT GTACCAGCTG CCCTGCGGCC ACCTCCTGTG CCGNCCCTGC
CTGGGTNAGA AGCAACGGTC CCTGCCCATG ACGTGCACAG

SEO ID NO:1841: (Length of Sequence = 332 Nucleotides)

GTGTGATTCC ATTTATATGA AATGTCCAGA ACAGGGAAAA CCTATTTNAG ACAACAGAGA CACAAAGTCG ATCAGCAGTT
GCCAGGGGAG GAGGAAGACG GGAGGGGAAA TGATTGCTTC ACGGGGTGAT GACAGAATGT TCCAGAACGT GACAGAGGTG
GTGCCTACAC AACTTTCTGG ATGTACTAAA TGCCGCTGAT TGTTCACTTT CAAGTGATTG ATTTTTAGGT TATTTGAATT
TCATCTCAAT TAAAAAACCC AAACACGCAA ACTGCTCCCG CCAGCTTCAG CCCCGAGGCA GACGGCGCAN CCGTGGGAGG
GATGCTGAGC CA

SEO ID NO:1842: (Length of Sequence = 246 Nucleotides)

GCTGGTCAAG GCAGAGTITTA CTGAACTNIN AGTTTCCTCC TGCACACACC GGGCATGACA CCTTCAAGTC TGNCCAGCAG
TGGGTCCAGA AAGTACCCTG TGTGCCTTGG ACGCAGAGGC TACAGTTCTN ACTGTGTGGC ATGGGAGCCT TCANAGTGCC
CTCGGGAGCT GCCCCTGGTC TTTGTCTGNA AAGGTGACTG GGAGGNTAGA AAAAGCAGCG GGCTGGCATT GTTTCGGGGG
TGGGGT

SEO ID NO:1843: (Length of Sequence = 313 Nucleotides)

ATTTATTGCA AACAAAATTG AGGTAAAAGA AGCTGACCCA GAACCCACGC CCGTCCAGGC TGGGGAAGTC TCTACTCGCC CCACACCAGG CCCCGAGCAC CGCGGGCCCG AAGCAGCCCC CAGAGGACAG ACGGGCCCTG CGCACTGAGG TAGCTGCATC TTAAGCCCCC ATGAGTACAA CTGCCCAGGG CTGCCCCAATT CCCAGAGGG AGGAGGAGAG AGAGGCAGGC AGGGGGAGCC CCGGCTTCAG GTGGGGCACA CCCCANACCC TCAACAAACC TTCCAGCCTC TTCGGGCTGG GGCACTTCCT GCC

SEO ID NO:1844: (Length of Sequence = 274 Nucleotides)

CTTCGCTTCT NAAAACCAAA CTCCAGCCGC TECCAGTCGG GACTTGGTCG CCCGNCGCTG CCAGAATGCT CCACTGCCAG CCGGCCCCCC TECCTCCGTT TCCCTTCTGT TTAGTGGCGA CACAGGCACC CAGCTTTIGGG GTGGTGGTGAACGTTCCCAGG GGTGCCAGGA GCCACTGGGA CAGGGTGAGG CTCCCAGACG TGCCCAGCTC TCCAGGGAGC TTCTGGNCCA AGGNCGTCTG AGGGATCTGC TCCTTAACCN CCCA

405

SEQ ID NO:1845: (Length of Sequence = 441 Nucleotides)

SEO ID NO:1846: (Length of Sequence = 255 Nucleotides)

ATGAATTCAT TGTGIATTTA TTATTCACAG TTAATCACTA CCTACCAAAT GCTATCCGCA GAGTTAAAGG ATTAAGTACA
TAGGICTTTA TTTAAACACT GATTTTTTTT TTTAAATATA TACACACAAA ACTTAGTTCA GCAAGGCTTC ATGATATACA
CCAATTCCAA AATAAAACAA TCAAATGGTC CNGGNGTAGA ATGCCAGATT CCTTTTATCA TCTGCGAGGA AAAGAGAAGC
AGGATGAGGA AGAGT

SEO ID NO:1847: (Length of Sequence = 3. :leotides)

CAGGGCACAC GCAGGACCAC TGTGGATTAG AAACCCAC:
GGAGCCAAAT CTCATTTGIN ACCCTCAGTC ACCACCCC:
GGAGCCAAAT CTCATTTGIN ACCCTCAGTC ACCACCCC:
GGAGCCACACCTC CAACATTCCT CCCACACCAC CATCCACGAC
GGACCACACCAC CACCATCCA CATCCACGAC
GGACCACACCAC CCCCAGGCTA GGGTGGGAGG ATTTAGAGCA
GGACCAC CCCCAGGCTA GGGTGGGAGG ATTTAGAGCA

GTGCAAGAAA CCAAGGAGGA TGGAGCATCC AAAGGAAGAL AGGCAGGC TNGGGGATTG AGGGCAGGAA GGGCT

SEQ ID NO:1848: (Length of Sequence = 311 Nucleotides)

CCACTGGCCT ACATTATAGA AGIGCTGTAT GCGGACCCTG CCATTGTCAT CATGGACGCA GGCCATGACC ATCATCACCA CCCATTTTNT TGTCTGAAGA GAATCCAACT GCTACCCCAC CATCTGTGTC TGCACTCAGC TCAAATTCTA CATCAGCCCC TATCATCCGG TAGCTGAGGA AATAGTCACA GGTCTCTGCA TTACAGCCTG GTTTGCCATA TCTAAAGCAT CCCTTAGTTT TTCCACAGGC GTCCACCTTG ATTTTGGCAA ATGGNTCCAC AGGAGAAGCA GCAGGGCTNN GTGTCGGGTG T

SEO ID NO:1849: (Length of Sequence = 318 Mucleotides)

GIGAGICCCC CAAGAGGGGC CTCAGTCAGG AATGICGATG ACCAGTGGGC ACAGGTGGAG TGAGTGCTIG ATGCCCATGG
TGAAAGCAGG GATGIGGGGC TTGIGCACAG TGANCTGCTG GACCTCGTGG GAGCCGGGGC CAGGCCGTGG CGTGAGGTCC
AGAGGGTAGG CGAAGGCTTG GCCATGCTGT AAGIAGGGCT GCGGTTCTNA TAGATGGATG GCTCAGGTCG GGCGTACGTG
GTAGGTCCAG GGCCTCCTGC CACATCCTCC TTGIAGANCC AGTTCTTGTC CCTGGAGGCC AGACTNTAGC AGGGAGCA

SEO ID NO:1850: (Length of Sequence = 406 Nucleotides)

GGAAGCCACT GATTITCCCT CCAGTATGAT GATTIACTIT AAAAATGAAC CCAGAGGGAC GGGCATGGTG GCTTATGCCT
CTAATCCCAG CACTTCAGGA GGCTGAGGCA GGCAGATCAC CTGAGGTCAG GAGTTCGAGA CCAGCCTGGC CAATATGGTG
AAACGCCTGT NICTACTGAA AATATAAAAA TTAGCCGGGT GTGGTGGTGT GCACCTGTAG TCCCAGCTAC TCAGGAGGCT
GAGGCAGGAG ACTCACTNAA CCCTCGTGGT GGAGGTCGCA ATGAGCCGAG ATTNCACCAC TGACTNCAGC TTTGGCAACA
GAGCAAAGAC INCGTCTTCA AAAAAAAATA ANAAGGGAAA AAAAACCCNG NAAAAGCTTT TTTATTGTTA AAAACAAGTG
GGTCAC

SEO ID NO:1851: (Length of Sequence = 328 Nucleotides)

CTGAGGGGCA TITTITATTA TAAATTTAAT ATGGTTGATT AATGAAAAAT GACAATGAAG TACCAAGAAA ATGTTTGTCA
ATATAAAAAT TITAGCAGCA TITCCATAGT TICAGGCTCC AACATTAGTC GTACTTCCTC CCTCCCGCTA TCAAAAAAAG
AAGAGACTCC AATGGGATGG AGTAGAGCCT GGGGGTGTCC AGCTTTGTGT GGGCCTCAGA GAAATACTCC ATCCAGCATC
CAGGATTCTC CCTCCCTCTC ATCCCTGAAG TGCTAGAATG TCAAAGCACA GAAAAAAGCCT CCTTTGTGCT GACATTGGAG
ACAAGGAT

SEO ID NO:1852: (Length of Sequence = 174 Nucleotides)

GGGCAGGACG GCTCTNGGCC CTTCCTGGCT GACTTCAACG GCTTCTCCCA CCTGGAGCTG AGAGGCCTGC ACACCTTTGC ACGGGACCTG GGGGAGAAGA TGGNGCTGGA GGTTCGTGTT CCTGGCACGA GGCCCCAGCG GCCTNCTGCT CTTACNAACG GAGCAGTAAG GACG

SEO ID NO:1853: (Length of Sequence = 252 Nucleotides)

GAGCCATGCA CACACACGGC CGCATAGTCA CACACGCATA TCTACATGTC CCCCCCACAT ATACACACAC ACATATACAT GGACCCATGC ACACACAGG CTGGATATTC ACACACACTT GCACATCCAC TCCATATACA TAGACACGCA CAGACACAGC TGCATGTTCA CACACGAGGA CGTGCACACG GACACAGACA TGCATGCATA TGCGCACAGG TGTGTACAGC CTCAGTGGTG GGGGTTGGCT GT

SEO ID NO:1854: (Length of Sequence = 288 Nucleotides)

GGAAGGAGG CTAAACAATG GTCTGCAGCT CAGTTACTCC TCATCCTCGC CTGGGCCGGG CCAGCATCCA CTCCCCTTCC
TGTAAAGCAT TTGGATTTCC TTGGGGAAAC AGCCCTGCCC TCTGTCCTGA TCCATGTGTT TTGAGATCTC ACAGTAGCAA
GTGACTCATG TTGGTTCAGT GATTCCCAGA GGCTGATTCA AGGATGTCCC CAGCTAGACC CAGGATGGTG GACTCCAGAT
TGGGGCACTG GGCAGTTTCA CATCCTCAAG GCTTGGCCAT CATCGGGG

SEO ID NO:1855: (Length of Sequence = 293 Nucleotides)

AAAATGCITG TIGATATITI AGITATTAAT TCATATTAAC TITGGCIGAA ACITITAAAT TCIATIGIGA ATAGICAAGI
AAAATTIAGA TIGITACATI CIGGGITAGI ATTAGATIGI TITTAAGAT GITITAAACA AGATGITITI AAGATGAGIT
TTAAATAGIT CICITAACAC AAATAAAGCI TAATATGAGI ATTIGAAGGA AATTATCCCA AACCATICCA GITCCIGGCT
GIGAAAGGCT TITCCAGGGC TAATAAGIIT TCCACITCAG CCGTAAGTAG GIG

SEO ID NO:1856: (Length of Sequence = 308 Nucleotides)

ATCTTAGCAG AATCTTGAAA AGCCCAGAGA TOCAAAGAGC CCTTCGAGCA CCACGCAAGA AGATCCATCG CAGAGTCCTA
AAGAAGAACC CACTGAAAAAA CTTGAGAATC ATGTTGAAGC TAAACCCATA TGCAAAGACC ATGCGCCGGA ACACCATTCT
TCGCCAGGCC AGGAATCACA AGCTCCGGGT GGATAAGGCA GCTNCTGCAG CAGCGGGCAC TTACAAGCCA AATCAGATGA
GAAGGCCGCC GTTGCAGGCA AGAAGCCTGT GGTAGGTAAG AAAGGAAAGA AGGCTGCTGT TGGTGTTA

SEQ ID NO:1857: (Length of Sequence = 299 Nucleotides)

GGGGAAAGCT AATTGGCAAT AATCCTTGGG GGAAGGTCAG ACTCCTCTC TACAGATCTA GGGAAGGCCT GGTAAAATGA
TGGCTCTTTG GAAAATGCCA AGCTCCTTCA GATTCCATAC CCTCTCGGGC CCTCAAGCAT AGGCAACGAA CTTGTTCCTG
GCTTCACGNT TTCTCATTGA ATCAAAGCTC TCATGCATGG CCTGGATTTG TAAACACATG CTGGCTGCCA GCAGTGGCAA
GTTAGCCTCC TGACCCACTT CTCTCCTCCT TTCACTCTGG TGTATGAAGG GGGATGAGG

SEO ID NO:1858: (Length of Sequence = 295 Nucleotides)

TAAGACITCC TGTTAGTAAA AGCTACCICA TGAAAAGTAT TGATGTTATT TGCCAACATT TAGACTAGCT TTTGTTACCG
TTTCAGTTAT TCAATTTAGT CAGCACATGT TTGAGTGTCT TACTGCAGGT GAATAATCCA TGATTTCTGC CCCAGAGTAG
TTCATAAGAC TGGTAGGATA CATAGATTTG TAAATAAATA ATTATAATTC TGGGCAGTAA GTGCTGCTAT AGAAGTCTCT
ATAAAGCAAT GTGCAAACAC AAGAAAAGGA GCCGTTAATT CCTTATAGGG AAAGG

SEO ID NO:1859: (Length of Sequence = 326 Mucleotides)

CTTTATTTAG TGCTGGGGCT TTGGAAGCAA ATGTACCTGA GTTTGAATCT CAGGGATAAC CTTTTGACTG TGGCCCTGGG
TAAGTTACTC ACTGTCTCTG AAACTTCAAG TTCCTCATAA ATAACCTAAG ATGGACAATC ATAACTCTCT CTTGGATTGA
GGTAGGAGAA TATGGTGGAG GCAGGGAACC GAAGGCCATT TCACTCCAAC TTCCTAGAAC TAAATTAAAA GGAAAACCCT
AATTTTCCAT GCCTAAGTAA CAAAAGGACC AAAGGTTACT CCGTTTGCAA ACTCCCACCT TTTCTGCATG GCAGATGGGA
AGTTGG

SEO ID NO:1860: (Length of Sequence = 294 Mucleotides)

CCACCCCTAA AAGCACCTGG CCCCGCTACA GCAAACCAGG TCTGTCCATG CGGCTGCTGG AATCAAAAAA AGGCCTCTCC
TTCTTTGCGT TTGAGCACAG TGAGGAGTAC CAGCAGGCTC AGCACAAGTT CCTGGTGGCC GTGGAGTCTA TGGAGCCGAA
CAACATCGTG GTTCTGCTCC AGACGAGCCC TTACCACGTT GACTCACTCC TGCAGCTCAG CGATGCCTGC CGCTTTCAAG
AGGATCAGGA GATGGCTCGA GACCTCGTAG AGAGAGCGCT GTACAGCATG GAAT

SEO ID NO:1861: (Length of Sequence = 183 Nucleotides)

SEO ID NO:1862: (1 th of Sequence = 296 Nucleotides)

TTCGGCTTCT TAAAGITC... CCCATCCCTC CIAAGGICTA AGATGATGCA TIAAACACAG AGGATGCCCA ACAGTGGCTG
ATGGAATTAC CAAGTAAAAT CIAAGAGGTA GAAAAATGTG GTAGTTTTTA AATTTTATTT TATTAGTATG CAGGTGGGAT
TCAGAGACGT AAGATCTTAG CCTTTATTTT CAACATCTCC CATGCATGTC AACAAAGATT ATCAAACACA GGAAGTGAAT
AAAATACTAT GTAGACACTG ACCCTCTTTA TATAAAATGT GATTGATCAG GTCTGG

SEQ ID NO:1863: (Length of Sequence = 259 Nucleotides)

CAAAACAAAA AGGGCTCAA ACCAACAGGA AGTCAGCCCC ACCGCAAGCC GGACTACAAC TAACTCGTGC TCTCCACGCT CAGGCGTGGA AGCCAAGGCT GTGCCAGGCC TGGCCAGGCC AAGCAGGATG ACAGCAAACG CATTCTGAAC GINTAGCAAT CAGGTCCCCT GTAATGTGCT TGGAGAGINT GGACAAGGGC CGAGATGACG AGCTATGAGC TGTGGAAGGG AATGGGGGAA GCAGAAGGGC ACAAACAGA

SEO ID NO:1864: (Length of Sequence = 290 Nucleotides)

ATCCTTACCA ACAATECTTC CCAACTECCT CAAAECTCTC CTAAATEAGA ACATAGTTCT TTCTGAGCAA GGTCCTGTOG
ACCATGAAGA ATGTCACCAA GCTCCCCTCA GAGTCAGCGG GAGCTCAGCC AAAGCACAAG TGCAGTGCCC AGCTCCTCCC
ACTCTGCACC TGCTGCCTCA NACTCCCCCAC GCTGAGCCCA GGCCCCTACC CTCTGAAGGT GTTTCCCATG TGATTCTGAC
ACACACACCCC CACAAGAACC AGATGATCTA TGACATACAG CATTTAGCTA

SEO ID NO:1865: (Length of Sequence = 236 Nucleotides)

CATTICTGIT ACATTGAGAC TICAGICACC AACATCIGGI GGCAGAGATA CAGGIGIATG AAACATTICT ATTTACCCAA
ATATGCCAGI TCCCAAATAG GATGACIGCA TITAGIGITA AACIGGCITT TCTCATTAGA TACTCIAATT GAGGAATATI
TAGCITCTIG AATAGAAACC ATCCAAATGA TGITTTITIT TIGATATGIC TGIAACTATA AAAATCAGCA AATAAG

SEO ID NO:1866: (Length of Sequence = 424 Nucleotides)

TACEGGAAGE CEGIGITIEG AGECTEGAGE CETEGCAACE TCATTGAGAA AATECTCACA GACCGCCEGT CTACAGACCT
TAATGAGAGE CECCGTECAG ACGTECTTEC CITCCCAAGE TCTGGCTTCA CTGACTTEGC AGAGATTGIN TCCCGGATTG
AGCCCCCCAC GAGCTATGIC TCTNATGGCT GTGCTGACGG AGAGGAGTCA GATTGTCTGA CAGAGTATGA GGAGGACGCC
GGACCCGACT GCTCGAGGGA TGAAGGGGGG TNCCCCGAGG GCGCAACCCA GCACTGCCTC CGAGATGGAG GAGGAGAAGT
CGATTCTCCG GCAACGACGC TGTCTGCCCC AGGAGCCGCC CGGCTCAGCC ACAGATGCCT NAGGACCTCG ACAAGGGTCA
CCCCTCCTCC ACCCTGGACT GGCT

SEO ID NO:1867: (Length of Sequence = 256 Nucleotides)

AAACAATIGA AATCCACAAG AAATTACTAA CAGCACGTGT TTACGTTTTA TCCTGAATCA TACATTTTAA CAATTCACAG CTACAGGAAA TCTAGAACAA AATCAAATAT TCATCACGTT GGGTTGAAAA GTTGGAAGAT TTTGCATCTT ATTGAAAAGA ATTTTTCAAA AATGTTTCTG TACAAATGAA TGGAATTGCA CCAGGCTGCC CATGGACACC AGGTGTGGCC GCTTCCCAAC GGTCACCCAC CAGCTT

SEO ID NO:1868: (Length of Sequence = 297 Nucleotides)

CAAGGITITI TITTATITGI AGCIATAGCI ACAACITGGC AGCATGGGG AGGGTGGGAA TGTCCTGGAG GGTCTCCCAG
CCCTCCGCAA GCAGAGTACA AAGGCTGCTC GGGGGCCCGG CCGAGGGCGC GGNTGCAGCA GTGNAAGCAG CAGCACTAAA
CCTGGTGCCC CCCTCAGGTG GGGTGTCTGG AAGACGGTGG GCAATCCCTG CAGGATGGGC GAGGACCAGA CCCCAGGGCG
GGGATCCTGC ATCCCTAGAC CATGTTGGGT CCTGGGTCAN GGCACCTNGG NATGCTA

SEO ID NO:1869: (Length of Sequence = 470 Nucleotides)

CAGACATCTG GAGCATGGGA CTGTCTCTGG TAGAGATGCC GGTTGGGAGG TATCCCATCC CTCCTCCAGA TGCCAAGGAG
CTGGAGCTGA TGTTTGGGTG CCAGGTGGAA GGAGATGCGG CTGAGACCCC ACCCAGGCCA AGGACCCCCG GGAGGCCCCT
TAGCTCATAC GGAATGGACA GCCGACCTCC CATGGCAATT TTTGAGTTGT TGGATTACAT AGTCAACGAG CCTCCTCCAA
ACTGCCCAGT GGAGTNTTCA NTCTGGAATT TCAAGATTTT NTGAATAAAT GCTTAATAAA AAACCCCCGC AGAGAGACA
GNTTTINAAG CAACTCATGG TTCATGCTTT TTATCAAGGG GATCTNGATG CTGAGGAAGT NNGATTTTTT CAAGGTTGGN
TCTGCTNCAC CATNGGGCTT TAACCAGNCC CGGNACAACC AACCCATGGN TGNTGGNGTT TAAGAGGTTTT

SEQ ID NO:1870: (Length of Sequence = 344 Nucleotides)

AGAGATTAGA TITGITAAAC ATCTAGGITA AAATGGITAA AAGGATTITC ATACAATTIT AGGCACTATA CACGITGITT
ACAACAGCAT TGGTACITGG ATATGGGGAA AGATAAATCC GACATTITAA TATCITGATC AATTTGTGAC ATTCAAAATA
ATTCCATTIA AGAAACATTA ATCAAAACTT AAAGAGACAT ACCACTAAGT ATCCCACACA GTATACIGAA AATAAATATA
GNAATACAAC CAGAAGTCTA CAGNICACCA CAGTAGACAG ACTGGTGAAG NCCCAGCTTT TCATGGGCAG TNAAGGGCTC
TGGGCTAGAT TIGGGTGTCA ACTG

SEO ID NO:1871: (Length of Sequence = 278 Nucleotides)

GGATTTATTG TCATTCCTCC AAGGTCAGCA GGGGAAGGGG ACACCAGCCA CACTTCACCA CAGGCATAGG TGGCACTGAG

CCACCTGGCA CTATCTCCAC GTGCTCCACA CGGAGGGGTG CCTTCTCACT GGCAGCAGCT GCACTTCTCT GCTTCTGCCT

CAGCIGCCIC TCCGCCITTG CACACACAGI CCITGGCACA CTTCTCACAC TNCGCAGGCA GCAGGAGCAG CAGCICTTCT TGCAGGAGGI GCATTTGCAT CCCTCGCACT TGCAGGAG

SEQ ID NO:1872: (Length of Sequence = 271 Nucleotides)

CTTGCCATCT TCACAGCCAG AAGCTTCCTT GCTTCATGCG CAGACCCTCG TGACTCCCCT TCCCTTATAA GGGCCCCCCAT
GATTACTCAG GGCCCACCTC AACCATCCAC GGTCATCTCC CCACCACGAA ATCCTGAACT GAAGCACAGG CGCCGGGTCC
CTTTTGCCAC GCAAGGTAAC ACTTTCCCAC GTCCTGGGGT TCCAAACCTG CACATCTCTG GGGCCTGTTA TINCACCCAC
CGTCATCAGT GAGGCGCCCTT NAGGAGGGGC T

SEO ID NO:1873: (Length of Sequence = 332 Mucleotides)

CAGGGTATAG TECAGTEGEG CAATCTCGGC CCACCACAGT CTCGACCTCA TEGGCTCAAG TEATCCTCCC ACCTCAGCCT CCCAAGTAGC TEGGACTACA GGCATCCTCC ACCACGCCA GCCAATTTTT TECATTTTTC ATAGAGAAGG GGCTTCACCA TEGCTCCCCA AAGTGTTAGG ATCACAGCCG CGAGCCCCTG GACCCCGCCT ATAGTTTTTG TTTCGCTTTG TTTTTGTTTT TTGAGATGGA GTCTCACCCT GTCANCCAGA TEGGGATGCA GC

SEO ID NO:1874: (Length of Sequence = 317 Nucleotides)

CTCTCCACCT CAACCTCCAG CCCACCTCCA GGCTGGGGAA GGGGCTGAGT CTTCCCCTCC CATACATACC TCACCCGGCC CCCAGCCCAC AGAGAGGCTG AGGGAGGGCG TCTGGGTCCT CCTCCATCCC TGTACCTGCT TCTTCCCTCT TCATTTCCAC CTTCTAGATC TTTCCCCCCA CCCAGCCCAC CTCCAGGCTG GGGAAGGTGA GGAATTCTTT CCTCCCACAC CCTACCCCAC CTCCACCTGCA GACCTGCAC GACCCACAC CCCGACCCT GCAGGCT

SEQ ID NO:1875: (Length of Sequence = 185 Nucleotides)

GIGITCCACC CACCTOGGCC TCCCAAAGIG CIGGGATTCC TGGCGIGAGC ACGCIGGGCC TGGACAGICI GCCCCTAGAT GAGITGCCCA GCACGGIACA GCIACIGCCI GCCCGGACCC CAGCCCCTGA TICIACOGCC GCICGGCAGG GGGACGGCCA GGGAGAGGIC CAGCCGCGCG GCAAG

SEO ID NO:1876: (Length of Sequence = 214 Nucleotides)

CCTGGGGACA AAATAGTCAG CAAATTCTCA AGGGGAGAAA ATAAAGTACT TCCCTTCTGT TAAAAAAAAG TCAAGAGACA AATCTTTCCT CCCCCATTCT CACTAATAGT TATTGAAGGG GAAAAAAAAA AACCCCACAA CTTTTTAAAC TAAAGATAAA AACAAATGAA AATGAATAAG ATCCAAAGAA TGTCTTTTGT TACTCTGCCT TATG

SEO ID NO:1877: (Length of Sequence = 340 Nucleotides)

TITICAACAAG AAGAAGITGA ATTIATCAGI GIGCCIGICC CAGAGITTGC AGATAGIGAT CCIGCCAACA TIGITCATGA
CITTAACAAG AAACITACAG CCIATTIAGA TCITAACCIG GNIAAGIGCI ATGIGATCCC TCIGAACACI TCCATIGITA
TGCCACCCAG AAACCIACIG GAGITACTIA TIAACATCAA GGCIGGAACC TATTIGCCIC AGICCIATCI GATTCATGAG
CACATGGITA TIACTGATCG CATTGAAAAC ATTGATCACC TGGGITTCIT TATTIATCGA CIGIGICAIG ACAAGGAAAC
TTACAAACIG CAACGGGAGG

SEO ID NO:1878: (Length of Sequence = 326 Nucleotides)

GAAAAACAAG GAAAATAGGC AACAACCTGC AATGGACACT TTTCTCTACA GAACCTTTTC AACCCTGAAT TGAATTGTTT
CCTATTCATT TNCTAATAAA AAGTTACTTT GCAAGATATA AGGAAATACT GTCCCAAAGA TTTTCACTAG TCATTCAATC
CATTAATAGG ATTTGAAAAG GCATCATTAC ACAGGGTTGA AAATACTCTG GAATGAGACT GCTTTACAGT CAGAATGCCT
GAGTTTTGAG GCACTGTTAC TTCTAAACAT CTCTAAGTTT CTATTTNCTC ATCTAAAGGA GTAATATTAC TTTCCTTAAA
AGGTTG

SEO ID NO:1879: (Length of Sequence = 222 Nucleotides)

GAAAGGGAGA GGTTGCAGCG AGCCAAGATC GTGCCACTGC ACTCCCACCT GGGTGACAGG GCAAGACTCC ATCTTAAAAA AGAAAACCCA GGAGTCTTTG GTTAATGTAG TGCAGGACTC TGAGCTCCCG GGAGGACCCT TCCCTCCCAG ATGAACTGTG ATGGACCAGC CCAAAGGAGG GGAGAGAGCA CTINGGCCAT AGTGGTGGTG GATCTTTCTA AC

SEO ID NO:1880: (Length of Sequence = 244 Nucleotides)

GACATGAATG GTATCCTCCT GGGGTATGAG ATCCGCTACT GGAAAGCTGG GGACAAGAA GCAGCTGCGG ACCGAGTGAG
GACAGCAGGG CTGGACACCA GTGCCCGAGT CAGCGGCCTG CATCCCAACA CCAAGTACCA TGTGACCGTG AGGGCCTACA
ACCNGGCTGG CACTNGGCCT GCCAGCCCTT CTGCCAACGN CACGACCATG TAAGCCCCCT CCGCGGCGAC CTCCTGGGCA
ACAT

SEO ID NO:1881: (Length of Sequence = 156 Nucleotides)

GTACAGGGA GAGTIGAGCI GIGACAAAGI CAAACACAGG CCTIGGCCAC CCACAGGAGC TCIGCAGCIG GGGIGGTCIT GAAAGITGIC TCAGIGAAGG CAAGGIGCIG AGCTIATTAC CCCAGCAGIC ATIGIATITA GGCICCGIGI GGIACC

SEO ID NO:1882: (Length of Sequence = 210 Nucleotides)

TTTTTTTGA AACGAAGTCT CAGTCTGTCA CCCAGGCTGG AGTGCAGTGG CACGATCCCG GCTCACTGCA ACCTCTGTNT CCCAGGCTCA AGCTAGTCTC CTGCCTCAGC TGCCCGAGCA GACGGGACTA CAGGCACCCC CACCACGCCC GGCCAATCTC CAAATGGTTC TTTTTTTCCG GAGTAGTAAG TTACAATATG GGAGATTATT

SEO ID NO:1883: (Length of Sequence = 214 Nucleotides)

GIGATGAATA CATCCAGITT TCCAACCACA TTCCACCAGG TGGGTGTTTG GCTGTGGGAC GCATTATGTA ATCTTCGTTG
CCAGGAAATT TACCTTCCTA ATTACATTTT GCAAATGTTC ATTTGAAGCC GCCTTCTTGG AGCTCACAGT AACTAGGAGG
TGGCTGCTGG AAGCCCCAGG GCACCGTGGG AGGGACAGGG GAACGTCCCA GACC

SEO ID NO:1884: (Length of Sequence = 211 Nucleotides)

ATCITICGCI CIATGIGCCA TCACCIGGAC ACTCIAGGIA ATACCCCCIG TIGGGCAGGG GIGAGCICCC AAGGCCICAG GCAACCCAGC TCCCATGACT TIGCIGGGCI CAGCCCACAT AACTGITCTC ACAGGATAGA GITGIACACT GGIGCITACA GCTITCCIGG GCCAGTGITG CATGCIGCCA GIGGCIGCAG CAGCAGCCCC A

SEO ID NO:1885: (Length of Sequence = 212 Nucleotides)

ATTACCIGAA TICCCGIGIG GCCGGITTGGG TAGGCAAAGG AGACATCITG GAACIGGACA AGGCCCTCCA AGIGTAAGGG
AGTCAACAGA CCACTGGGIG GGCAGCGAGG GGIGCCGTCC AGGTACTCAA ATATITICTC TGAGGAGCCC ACAGCCTTCT
GTACTCIGGG GTAGATGGAG AGCAGTACCT CCACAGCCIG GGIGAACIGC AT

SEQ ID NO:1886: (Length of Sequence = 208 Nucleotides)

CATCCGCATA GIATTIACAT CATGGGTATA GGCAAGINCI ACAAATCAGG NCTTINCCIT GGGGATGGAT GITTGGAGCI AGITTACCAG CACACCAGTG GGTAAAAGTG AACAAATACT TTTTTGATCC CACAGAATCT TAAAAAATAC TTTACTTCGA AAATGTCTCT ACTAAGTAAT CATATATATA TATATATNIG TATATATA

SEQ ID NO:1887: (Length of Sequence = 332 Nucleotides)

CTOTATACA COCCECAR TOCCATICA ACTICCTITI TACACTEGAT GITICIATCA CATCCIGAGE ACCACIAACC
CACCAGCAAG TCTCCCCCTE ACACACATIC ACGIAGGICC ATACCCTICA GAGICCIAAA GGGITAATGA GAAGCCACCT
CAGCTITIGGI GAATGGAGCC CCAGCCCCAA ATCCCCTCCC CTTGCAAATA TGGGACAAGI AGGGAGAGIC TGATGGAGGC
ACCAGGACAA CTACAACAAC CTCTTACCCC TCAGCTATAG ACACCTAGAT CAGGACAGAG GGATGCATAT GCCCTCTCCA
CCTTAACACC AA

SEO ID NO:1888: (Length of Sequence = 224 Nucleotides)

AAGAGCTGAT TGAGGCTGCC AAGAGGAACG ACTTCTGTAA GCTCCAGGAG CTGCACCGAG CTGGGGGGGA CCTCATGCAC CGAGACGAGC AGAGTCGCAC GCTCCTGCAC CACGCAGTCA GCACTGGCAG CAAGGATGTG GTCCGCTACC TGCTGGACCA CGCCCCCCCA GAGATCCTTG ATGCGGTGGA GGAAAACGGG GAGACCTGTT TACACCAAGC AGCG

SEO ID NO:1889: (Length of Sequence = 261 Nucleotides)

CACTITACIG AGTCACACCC AGCTGIAAAC ATGTCACCGT GAGANTCCCG CCCCCCACCC CCAGGCCGCA CAGTCCGCGA TGAAATGACA GGGGAGCGGG GAGGGTCGCC GGAGCGGGTG CCAAGCAAGG CAGGGCAGGC AAGTGCAGCA GGCCCTGAGT TTCCGGGAGG AAGCCCGGAG GAGGTGGGGT GGGGCAGGAG CGNGGGCTGG GGACCCGGCC GAAGACCAGG GGGCCCAGGA AGCCTCTTTT CCGAAGGNCT T

SEO ID NO:1890: (Length of Sequence = 312 Nucleotides)

CTGCGAGACT ACGAGACGGT GGTCAAGGTG AAGCCCCATG ACAAGGATGC CAAAATGAAA TACCAGGAGT GCAACAAGAT
CGTGAAGCAG AAGGCCTTTG AGCGGGCCAT CGCGGGGGGAC GAGCACAAGC GCTCCGTGGT GGACTCGCTG GACATCGAGA
GCATGACCAT TGAGGATGAG TACAGCGGAC CCAAGCTTGA AGACGGCAAA GTGACAATCA GTTTCATGAA GGAGCTCATG
CAGTGGTACA AGGNCCAGAA GAAACTGCAC CGGAAATGTG CCTACCAGAC AGAGAAGATT ACAGTATGTG GG

SEO ID NO:1891: (Length of Sequence = 298 Nucleotides)

CCTAAAGGCC AGGCAAGGCT GATTCTCCAC TTCCACATGA GACAGAGCTG ATTCTGCAGG GAAACGGCTG GGGAGGCTCC ACCTCTTTCC TCCCCACAAC CATTTACTGG GAAGTTGTGT ATACTTGGCA GINTGGGAGG AAGGTACTTG GAAGACCCTG CCAGCCATCT CCCACCCAGA CTTCTTCTCA CCAGCACAGT CTTCAAGGCT TGGTGGGAAA GGTGTGTGGG AGTGGAGAAA GACAAAGGGC CCTTCTTNAA GAGAGGAGCT GCAGAGAGGG GCAAAGGGGT TCCTAGCC

SEO ID NO:1892: (Length of Sequence = 333 Nucleotides)

CICCAAGGIC ATCCAGICCG TCGCTAATTA TGCAAAGGGI GACCIGGACA TATCTTACAT CACATCCAGA ATTGCAGIGA
TGICATICCC AGCAGAAGGI GIGGAGICAG CGCTCAAAAA CAACATCGAA GATIGCGGIT GITCCIGGAC TCCAAGCACC
CAGGGCACIA TGCCGICTAC AACCIGICCC CGAGGACCTA CCGGCCCTCC AGGITCCACA ACCGGTCTC CGAGIGIGGC
TGGGCAGCAC GGCGGGCCCC ACACCIGCAC ACCCIGIACA ACATCIGCAG GAACATGCAC GNCTGGCTGC GGCAGGACCA
CAAGAACGIC TIC

SEO ID NO:1893: (Length of Sequence = 487 Nucleotides)

CCAGATAGAG TITCTGTTT TAAGTTTAC ACGTGCCACA TCAGGGAAAG TTAGGTTATG ATTAAAGCAA GAGATGATAG ATGAACAAAC AAAGAAACAA CAACAAAAAG CCCATGCAAG AGGCAGGAAA AGAGGCTGAC TGGTTAAAGA ACAGGCCAGA TTGGACAATA CTGATCAAGA GGGGTTCACA TTTGAAAGAA CAGTGCTTTA TTCCTCTACT GACTAGAACT AAAGGGATTT TGGCCGGGTA CGGTGGCTCA CACCTGTAAT CCCAACACTC TGGGAAGCCA AGGTGGGCGG GTCACGAGGT CAGGAGTTCG AGACCAGCCT NACCAACATG GGTGAAAACCC CATCTCTACC CAAAATACAA AAACTTTTNC CGAGCGTGGG CCCGGCGTTG GTTGGCTCAT ACATTTNATN CCCCCNCTTT NGGGGGCCCA NCCGGGGGGT TCACCTTAGG GTCAAAGGGT NCGGGGNCCT TCTTGGC

SEO ID NO:1894: (Length of Sequence = 283 Nucleotides)

GETGETGAGE TEGECTICTEG AGAAGCTEGA GCTGACCAAG TACGCAGACA AGCCGGCTGG CACCTACAGC GGCGCAACA
AGCGGAAGCT CTCCACGGCC ATCGCCCTCA TTGGGTACCC AGCCTTCATC TTCCTGGACG AGCCCACCAC AGGCATGGAC
CCCAAGGCCC GGCGCTTCCT CTGGAACCTC ATCCTCGACC TCATCAAGAC AGGGCGTTCA GTGGTGCTGA CATCACACAG
CATGGAGGAG TGCGAGGCGC TGTGCACGCG GCTGGCCATC ATG

SEO ID NO:1895: (Length of Sequence = 234 Nucleotides)

ATGTCCATTA GCCTCATTTG TCATCTGAGG GAGCTGGTGA GAACAGCCTT GGCGTGAAGG CATCCCTGGT AGAAGTCGGG GGAGATAGAT AGTCACAGTT CCCCAGTTGG TGGAAATNGG ATNGGAGTAG GGAGAGGCTN GAACAGACCC TTCCCCATTC ACCTGGNGAA TTTTCTCCTC CCACTGCCCT AAACACTTTA TTTCCATCAC AGGGGAGAAA TNCTGCTGAG AAGG

SEO ID NO:1896: (Length of Sequence = 285 Nucleotides)

CTTTAAAGIG TAATAATATG ATTTTTTAAA AGAAATTTAT TACTIGITGC AAAGGICTIT TTAAACCAGI TTAGATTICA AGAAAAAAAA AATGGAAATC ATCGAAAATT CATTTCACAT TAATGGICTA AAAATAAACC AAAGGACATT ATGIGIGCAT GIGIGIATAA GIGCACACAG AAATATATAT NCATATGING ACTATATACA TGIGIGIATAA TATGIGIATAA TATACATNCA CTTGTATAAAA TGIATATACA CATATACCTA TAATGIGIGI ATGIG

SEO ID NO:1897: (Length of Sequence = 288 Nucleotides)

GCAGGITIAT GITITIATIT ATGIATINA ACTGACITAT TIGIGIATCC CACTAGAACA ATACATICAC AATATACITG CAGAACTGG CCTGGGGGAT CATGGGAGCA GAGAACTTGI CCAGIGAATA GITGITGAAG AAAGGAGTAA AAWCTCCCCC AAACCCTAAA GGCATCCTTT TCGIAGIGIG TGICCCAYAG GITATGGCTGC TGAGCACCAG GGGCTGCTCA CCATGNTCCC AAGAACCAGA GTCANGGAGG CAGACAGCAG GGKTTATTAA GGTGCACA

SEO ID NO:1898: (Length of Sequence = 398 Nucleotides)

CAGAAGTAAA AGATTITTAT TGITCTATAG ACACTTCTGA AAAGAGATCT AATTGAGAAA ATATACAAAG CATTTAAGAG
TTTCATCCCC AGAGACTGAC TGAAGGCGTT ACAGCCCTCC TCTCCAAGGC TCAGGGCTGA GAACGGTTAG CATATCGAAT
GATCAGTAAA AACATGCAAA AGTGAGAAGG AAAGGGAAAA AGGTGCATTC CCCTAAGCTG AGGGGGATGG AATTTCAGAA
CAGAGGAGGC AGGGTGGACA AGTACCAGGT GGCTCTCCCT TTCCCTCTGT GTTATCTTTC AAAACAGTTC CAAGCTTTGA
GAAAGCAATG AGCTCCACCT ACTCAGCAGA CCCACGGGTC GTCCCCCTGG ACGTGACTTA GCAGTGACCT TGCCTGCC

SEO ID NO:1899: (Length of Sequence = 227 Nucleotides)

CATGGGGACC CGGGTTTATT TTATTAGGAA GGAAACAACC AAGCACCCCA TGTTCCTGCC CGGAACTCCC GGGGGGAACA
TGCCAAAMAG CCGGGGATCG AACCCAGCCC ACCTGTCGTG GRGGKCCCTT CCTTCTCAGG CCACAGAAAT AAACCCGTGT

ACTITYTIATI GITAGCACAA CATTACCAGA AAACEKTAAC GGCAGCCAAG CAGGACAGAC AGTTAAG

SEO ID NO:1900: (Length of Sequence = 405 Nucleotides)

GGGATGCACT GGGITTCACA TCAAGTICIT GAGAGGWICC CGAACGACTT CTCTGCCCCA GGGGAGTCCG AGCCACAGTT

TTCTGATCAA CTGATGATTC TRACCCGCTT CTTTCTCTCT GGGGGGTAAG ACACTTGTTG TTGAGCTCTG GGGATGATGG

AGAACGACTC CTCGGCCTAG GAGTCTGAGG CAAAGCTTTC GGTTCTGGGG AAGAATCACA TTCGCTTCTC CCTCTAGATG

GCGTTCTAGG TATATCTTTC ATTCCAGGAG AGGACCCAGA CAGGCTGTGC CTCGAGGGAG TCCCAGACCC ATCTCTAAGT

CCTGGAGGAAG ACCCAGACCT GCTTCTCCTT GATGGAGTTC TGGTAAACCA TCTTTCATTT CAGGAGAAGA TGCAGACTAC

TTCTT

SEO ID NO:1901: (Length of Sequence = 244 Nucleotides)

ATRATICATA TECTAGITTA TITIATCITAT TATIGAGAGA TAATITCATG ATGACAGITA TCAATAATCA ATTACAATAT CAAGAAATIC AAAGAACAAA ATCITGCAGA GACTATGCTT TIGTATITGG ATTTAAAAAG TATGIGATCT CATTITCACA TACCAAGCTG AGAGGCCATT TAGACTATCT CITIGCTAAT TITIGCTTAC TGCTGTAGGG AAGAAGATTT CCAATGAMCT TIAG

SEO ID NO:1902: (Length of Sequence = 329 Nucleotides)

TAAAAATAAA AAAATAAATA AAATTTIAAA AATAATAAA ATTCACTATA TACACATATA AAGAAATAAA AAGAAGTCTC
AGTTGCAGCT ATTTGTCAAA ATTAATATCC ATTTCTWITW ATATACOGTG AATATTGCGC AATTATAGAT CTGGATTTTA
AACCACTIAA TGAAGCGGCA ACACCAGGTG TTTTAAGGTG TTGGCATTCT TCGCTGATTT GGCTGTTCCC AATGTTTACA
TTATTTAATC TTGCAAAAAT GGTTCTGATG CACTTGGGAT GTGAAATGCT GTCCCGTTTT ATTTTTTTAA TGTTGTTATC
CTTGGGTGT

SEQ ID NO:1903: (Length of Sequence = 421 Nucleotides)

ATTITATATI CCACAGICAG GIGGGICIGC GATASTCATI TAATGITAAA CGCCATCAGG GGCCICTCCT CCCGITTCTG
CCAGGGGCTT TICTIGICIT CTCCTIGGIC ATCATCATCA TOGICTICCT CTTCCTCGIG GGCAGATCTT CTCTGGIGGG
GGCTGGCTGC TGGCTCCGAG GGGGCATCCG CAGTCCGTCT GGTCGTCTCC TCCTGCAGGC TGGGCAGCTG GCCACCACTT
CTCCGACTCG ACCCTCCAA CAAGCATCGC AGGGCACTGT CCTCGGGGGT ACAGACCGTG GTCCCACACTT CGCTACCACT
CTGTTCCACG NCATCCAGGG TACACGAGCT GCGTGTAGGC CGTGCTGTCT TGGGGCTCGA GGCTCTTTCT GCTGGTGCTC
TTGGACGGGC GGGTAAATTC T

SEO ID NO:1904: (Length of Sequence = 423 Nucleotides)

GICTIGICGGC CCTGICTGAA GTGACGGTGC AGCCAGGCTG CTCCCTGCCC AGCCACCCCG AAGCCATTGT GCTGGACGTC
GACTACAAGT NIGGGACCCC GATGCAGAGT GCTGCAAAAG CCCCATATCT GGCCAAGTTC AAGGTGAAGC GATGTGGAGT
TAGTGAACTT GAAAAAGAAG GTCTGCGGTG CCGCTCAGAC TCTGAGGATG AGTGCAGCAC GCAGGAGGCC GACGGCAGAA
GATCTCCTGG CAGGCAGCCA TCTTCAAACT GGGAGACGAC TTCCCGGCAGG ACATGCTGGC CCTGCAGATC ATCGACCTCT
TTCAAGAACA TCTTCCAGCT TGTCGGCCTG GACCTCTTTG TTTTTCCCTA CCGCGTGGTG GCCACTGCCC CTGGGTTCGG
GGTGATCGAG TGCATCCCCG ACT

SEO ID NO:1905: (Length of Sequence = 370 Nucleotides)

CAGAACCAGA ACATTITIAC TCTTTGGGCT CTGGGAAGGG CCAGGCAGAG TGCAAGGTGT CCACAGGAGG GGTAAGCAGA
GAGGAGCTAC AGGGGGGTGC AGTCCTAGTA CCCTGTTGGG GAGGACTGAG GGATGGTGAG TTTGGTCTCC GGAGGGGGGCT

THE STATE OF FIRE

CCAGTCCTGG TGCCCAGTTC TNACANCTGC CCCTCCTGAG TTCACACTGG AGTCCTTGCA GTCCTGAAAC CACAAGGCCT NCCTGAACCC TGGGTCAGGA GAGAAANACT TGGGGAGGG AAAGGACGC GTGGGCTACC CATKACGGCT CTGAGTTCTT CCTGGGGCTT GTGTCTTTTC CTTGGCAGAA GAGGGCACAG CCAAAGGCAA

SEO ID NO:1906: (Length of Sequence = 415 Nucleotides)

GICACACCIT CATTCAGIGA GGAAGAAATG CTITCACTCT GGGAATTCAC AGCATCCCAA TCTGACGITG TACCCGIGIG ACACTGITTG TGAGCCCCAA GITTCAACGA GCTCITGCAA GTAAACGGAC ATTCGICACA TTTGIAGACA GCTGTCTTTC CAGATAAGIG GATGITTTCT ATGIGACGAG AGATGCTACG TCGATGCATG GTGAGGAAAG GACAAAAGGG GCACTGGAAC CTATTCATGA ATCINCTAAA TGGAATCCCC TTGGICTCCA ATAATTTGIT GCCATCTGAG CCCATCAGCT GCTCTGCAGA CAGGCCTGAT GTCTGGTGAT CCACAGCACT TAAACCATTC TCACTTGTCT ATTTCATTTA ACTCTTCATC AGAACTAGAG TCATTAGCAT GCTGT

SEQ ID NO:1907: (Length of Sequence = 214 Nucleotides)

TGAAATCCTG TACGTGTCAA CTTTGAAATG TATGTGTGT GGTTGGGTGG TGGTGATGTG ATACGGTTTG GATGTCTGTC
CCCTCCAAAT CTCATGTTGA ACTATAATCC CCAATGTTCC AGTTGACGTG GTGTTTGGTT CCATGGCGGG GTACCCTAGG
GATTCATCTG TTTTCTTCAC TTCCCTTTGC ATCTGAGATC CTGCTGGAAA CCAC

SEO ID NO:1908: (Length of Sequence = 410 Nucleotides)

CAGGAGAGCT GGGCACATGT CCCAAGCCTG TNAGTGGCCC TCCCTGGTGC ACTGTCCCCG AAACCCCTGC TTGGGAAGGG
AAGCTGTCGG GTGGGCTAGG ACTGACCCTT GTGGTGTTTT TTTGGGTGGT GGCTGGAAAC AGCCCTCTCC CACGTGGCAG
AGGCTCAGCC TGGCTCCCTT CCCTGGAGCG GCAGGGCGTG ACGCCCACAG GGTCTGCCCG CTGCACGTTC TGCCAAGGTG
GTGGTGGCGG GCGGTAGGG GTGTGGGGG CGTCTTCCTC CTGTNTCTTT CCTTTCACCC TAGCCTGACT GGAAGCAGAA
AATGACCAAA TCAGTATTTT TTTTAATGAA ATATTATTGC TGGAGGCGTN CCAGGCAAAG CCTGGCTGTA GTAGCGAGTG
ATCTGCGGGG

SEQ ID NO:1909: (Length of Sequence = 339 Nucleotides)

AAAATTAAAT CCAAATTITA ITAAGGATIT CAGGITACAT ACTICAAATT TCIAGAATGG AATGGAATCA TITTGGAACT
GGAAAAATGG CATAAACACT GACGICCCIT AAAACTICAA TITTATAAAG AAAATTCITC TGCAAACCAC ATCCCCITTA
TGIAACAAGA CTAGGIATTA TCIACACCIT CACITTGGCA ATAGCIATTT CCTAAAGAAT GAAAAAGATG ATTITNCTAC
TTCAGITCAT TAAAAATGGG ATTCTATCIT TGAAGITCAG AAAAAGCIGC ATTTCGATGA ACTATGGGTT AAAAAAAAAAA
GCACATAGIG TCTAATCAA

<u>SEQ ID NO:1910:</u> (Length of Sequence = 439 Nucleotides)

GECCCAGGGA GCACCAATCA CAGCAGGGGC TCTGGCCCAG GTGTCGGCAG CCCAGGCCTC CATTTGCTAA TGATTAATAC
ACTGTTTGGG CTGGCCAGTT TTTCATGCAT GCAGCTTGAC GATTGAGCAC AGTCAGGCCT TTGTATTAAA AATGAAAAAAT
GAAAAAAACAA ATTCAAAACC TATTCAAAATG GGTTCTAGTT CAATTTGTTT AGTATAAATT GTCATAGCTG GTTTACTGAA
AACAAACACA TTTAAAATTG GTTTACCTCA GGATGACGTG CAGAAAAATG GGTGAAGGAT AAACCGTTGA GACGTGGCCC
CACTGGTAGG ATGGTCCTCT TGTACTTCGT GTGCTCCGAC CCATGGTGAC GATGACACAC CCTGGTGGGC ATGCCCGTGT
ATGTTGGTTT AGCGTTGTCT GCATTGTCTA GGAGTGAAC

SEO ID NO:1911: (Length of Sequence = 342 Nucleotides)

AATGCACCCA TTTGGCTGCC AAGAGCTTCT CACTGCCTTG CTAGCAGCCT GCCACTGTNC CCTGGCAAAT TGAAACCACC CACGCAAACA CTCAAAACCC CAATCTCCTT GCTAATAAGA TACAACCAGT TAACACCGTG AAAAATGCAC ATCTCCAGCC TTCATTTCAA AAAAGAGCTC TGTACTAAAT GCAATATGCT TTTAAAGGGG GTTTTACAGG GACCAATCTC AATGCAAAGA CCAGTACCAG ATGTCTGAGT TTTGGTTACA GGTTTATAAAT TAGACACAAA ATTCACTCCA CACTGGAGTT TTACTTTCAA GCTGGAGTTA GCATTAGTTC TA

SEO ID NO:1912: (Length of Sequence = 380 Mucleotides)

TCATGCTTT AATACAAACT TAAAAAAATC TGGAACAATA GAAACTGTAC AGATTTGATC AATCTTTTIG TTTTGTTTTT
AAACTAAAAT CTCTAAACAC ACCAATGTCC CATTCCAAAA TATTGCACAA CATTCTGAAT ACAAAACCCT TGATTGTATT
CCTCCINCAC TAAAGAAAAA AGTTCATGAC CCTGCTCCCC GGGCTCCTCT CCAGGCTTGC CTCAATGCCC CCTTCCCATC
CCTAGGGAGA AAACTAGAGA ATCTATAACT CACTGCATTG AGAAAAACAC ATCATTCTGG ACTAACAGTT TTCCATTCTT
CAGANGGNIA ATCCACCTTT TGGATTTGTT CCTGGGGAAA GAGGGGTAGA TAGAGGGATG

SEO ID NO:1913: (Length of Sequence = 361 Nucleotides)

GAGACAGAGT TITECTCGIT GCCCAGGCTG GAGTGCAATG GCGTGATCTC AGCTCACCAC AACCTCCACC TCCGGGGTTC
AAGCCATTCT CCTGCCTCCG ACTCCCGAGT AGCTGAGATT ACAGGCATGT GCCACCACGC CCAGCTAAGG CTTTGTATTT
TNAGCAGAGA TGGGGTTCA CCATGTTGGC CCGGCTGGTC TCAAACTCCT GACATCACAT GATCCCCCCG NCTCAGCCTC
CCCAAGTGCT GGGATTACCG GTGTGAGCCA CTGCCCTGGG CTCTCCAGTA CATTTTTAGG GGGACGATCA ATGAGGATTC
TCTTCTCTGA GTTACTGCAT GTGTTACAGT TTATAATCCT T

SEO ID NO:1914: (Length of Sequence = 409 Nucleotides)

GGGGGCCTTA CAACTAGGTA TGGTGGATAT TGCCCGACAG ACGGTTGAAT TTCTCTACGA AGAGAATGGT GGCATCCCAA
GAGACCTTTA TCTTCCCACC ATTGAAGACA TTAAAGACGA AGCAAACAAG TTCACAATTG ATAAAGTTCG AAAAGGTCTC
ACAGTAGTAA CCCGCTCTCC AGACAGCAAT AATGTAGCCA GCAGTGCTGT TGGAACTGCT CTGCCAAAAAT TTGCCATCCG
AGGGATGCTG AAAACCTTTG GGCTTCATGG AGTCGTCTTA GATGTTGATT CAGTGAATGA ACTGGTGCAG GTAGAAACGT
ACCTCCGCAG TGAAGGTGTG CTGGTGCGAT ACTTGGTATC CTATTTGACA TGTTGGGAAA GGGCCCCCAG CAGGCTACCG
AARGGACTT

SEO ID NO:1915: (Length of Sequence = 402 Nucleotides)

ATGGITTATA GCAGGAATAC TIGITCIGAA TGACTIGGAG GGAAAGIGIG TGIGIATATG TGIGIGIGIG TGITTGITAG
TTTTTGIGAG GIAGGGAGA CTATTTTGI GGITCAGTCA CTCCAATTAT TGCCACAATG CACTITCCIT CATAACIGCC
CCACCAAAGG TCTTAAAAGC CATTTTTGGA GCCTATIGCA CTGTGTTCTC CTACTGCAAA TATTTTCATA TGGGAGGATG
GITTCTCTT CATGIAAGIC CTTGGAATTG ATTCTAAGGT GATGITCTTA GCACTTTAAT TCCTGTCAAA TTTTTTTTGGT
CTCCCCTTCT GCCATCTTAA ATGGTAAGCT GAAACCTGGG NCTACTGTGG CTCTAGGGGG TAAGCCCAAA AGGCCAAAAA
AA

SEO ID NO:1916: (Length of Sequence = 382 Nucleotides)

GARATGAGAC TITATICIGA AATTATIAAA AAGAACAGAG ATGCICCATT IGGCIGCATG CAGGGGGGG GGITGGGGGG ACAGAGGGGA GGACAGGGGC TCAGCCAGGG GGACCGIGIC TCITILCCAC GCAGGACACT GIGCATGGGG CICTGGGIGC ATCIGCCCAT CIGICIATUG GCCUGIGIGI GIGIILAGAGC CCAAACACAG ACAGCIGGGG GGCIAGIATC CIGGAAGGIT TCACTIGGIG GCITGGCCTA

GGGACCAGCA AGGGCITGEN GTTGGAAGGG GTGGCTCAAG GAAGCCTCTT TCTCCACTCA CA

SEO ID NO:1917: (Length of Sequence = 375 Nucleotides)

GAGATTAAAA TAAACAACAC AAAATGTATT TAAATGAGAA ATTGAAATAT TAAAAATAAT ATTAGGTGAC ATTAAAACTG
TCATAGAAAT AAACTGTATA TACAACAAAT AAATCAATGA TTGTTAACTT TTTTAGACAG TTTGAAATATC AGATTATAAT
GAATAGCATT ATTAGCCAGT AAAAAAGAGCA TATAAAATTAT TTTAAAAATTC CAAATAAAAA TATTTAAAAT TTTGAAATTT
TGGACCCAAA ATTATGTCAG TAATTTCATG AAAGTAGATC TCCAATAGGT CCTATATTCT AGACACTATG AAATGACATC
AGAAACCGTC AATTAAAGTG TACCCCACAA GTGATAACTA GCTACCATAC AAGTT

SEO ID NO:1918: (Length of Sequence = 315 Nucleotides)

AATATACAGT ATGATACACT GATGIGCAGA ATGIGATTAG TITATTAATC ATATGIGAAA ATATTAGIAG CTACATATGG CCAGAATAGA TITTYCTCTC TACAAATGIA AGITAGIGIT GATAGAATTT GITATGCGAT ATTIGGITCT TIGGITTCAG TCCAAATGCT TTCTTCTIGG CATTTCATTG ACCCTGIAAA TTAACCTCAG CATCAATTTT CITTTAAATT CAACAGITAT TCAAAATGAT CGGAAATTAA ACTIGIATGT AGCTAGITAT CACTTTGGGG GTACACTTTA ATTGACGGG TTCTG

SEO ID NO:1919: (Length of Sequence = 285 Nucleotides)

CAGAAGTAAA AGATTITTAT TGITCTATAG ACACTTCIGA AAAGAGATCT AATTGAGAAA ATATACAAAG CATTTAAGAG
TTTCATCCCC AGAGACTGAC TGAAGGCGTT ACAGCCCTCC TCTCCAAGGC TCAGGGCTGA GAACGGTTAG CATATCGAAT
GATCAGTAAA AACATGCAAA AGTGCAAGA AAGGGAAAA AGGTGCATTC CCCTAAGCTG AGGGGATGG AATTTCAGAA
CAGAGGWGGC AGGGTGGACA AGTACCAGGT GGCTCTCCCT TTCCC

SEO ID NO:1920: (Length of Sequence = 181 Nucleotides)

GCAGGITTAT TITTITATITI ATGTATITNA ACTGACITAT TIKTGTATCC CACTAGAACA ATACATTCAC AATATACTIG CAGAACTKTG CCTGGSGCAT CAGGGGAGCA GAGAACTTIT CCAGTGAATA GTTTTTGAAG AAAGGAGTAA AATCTCCCCC AAACCCTAAA GGCATCCTIT T.

SEQ ID NO:1921: (Length of Sequence = 351 Nucleotides)

AGACGGGGTC TCACTCTKTC GCCCAGGCTG GAGTGCAGTG GCGCAATCTC AGCTCACCGC AACCTCCGCC TCCCAGGTTA
AAACGACTCT MATGCCTCAG GCTCCCGAGC AGCTGGGACC ACAGGCACAT GCCATCATGC CCGGCCAACC TTCTGTACTT
TWAGTAGAG ACGGGGTTTT ACTGTGCCAC ACAGGCTGGT CCCGAACTCC CGACCTCAGG CGATCAGCTR CCTCAGCCTC
TCAAAGTGCT GGGATCACAG ACGTAAACCA CCATGCGGGG CCCCAGTCTT TTCTTCAGAG GGCTCCTNAG CACCCCCAAC
CCCAAACCTG AGGCCTGTGA GAGTCTATCC G

SEO ID NO:1922: (Length of Sequence = 198 Nucleotides)

CCTCATCTGG ACACAGATGA TITIGCCAAAG AAGCGGCCTG CCCAGATCTG CAAACCTTGC AACCCAGCAC TCTTGCATAT CTCGCTTAGC GTGTCCACAA CTGGGATGCT AGCTGGCGTA AAGATGCTCA CGCAGCCACC AGTGCCTCTG CCGTCCATAA GTGCAGTGTG ACTTACCCTC TGAGAGTGGC ATCTGCTG

SEO ID NO:1923: (Length of Sequence = 303 Nucleotides)

TIGATITICC TATGGTGTGA AATCCTTTGT TATTTTTCTA AAAAAATAAA ATTTAAAAAG AAAGAAAACT AAGGAAGAAC
AAGANGCTAT TTACCCAAAG TGAGCTINCA GITTTAGITT TGCATGGCTG TTTGACTGCC TTTCCGCCCT ATGAAAATCA
AGAAAATCTT TTTTAAAAAT GGAGTCCTGC TATTTTCCAC TCCTTGCAGA TAATACAAAT TCAGTTTGTC AGGTTGGATG

GTGAGTTGGG AGCTGTGATG GATCTGTTGG CGGGTTTTGG ATGTGTAAAG AATGATATAT ATA

SEO ID NO:1924: (Length of Sequence = 231 Nucleotides)

GICCICCTG ATTCICAACC TITGCAACCI GCCITCCGIC ACTGCIAGGI CCACGIAGGC TIAACCITGA TCITATATGI AGGACCGGIC TICACCITAA GCAAGAGAA TGIAAGAAGI GNITICCCAA CICAGIIGCI GGCCCAGCII TGGCCICGIG TICCCITICI GAGGACTGAC CITIGGIATI GCICIGGAGI CICATATCCC CITIGGCCCI AACTGACCAC G

SEO ID NO:1925: (Length of Sequence = 249 Nucleotides)

GITTTIACIT AACCATTCIA TIGITGGGAA TIGGGITICC ACTITITINI TATAGATAGI GGIGCAGIGA ACATITITAA ATAGCITITI NCITCAGIGI AATTATTICC NI'AGAGAAAG TIACCAAGAG TGGITTIACI AGITCAGAGG GCITCAGGAT TI'NATGGCT CI'NCI'AGCG GIGCICI'ATT ATCCINNAGA AGACITGIAT TACITCCAGT GICAAGAAGG TIGCNCITCC ATGGAATGG

SEO ID NO:1926: (Length of Sequence = 367 Nucleotides)

TTTTTCICAG CAAGGAACAG TCATGAGAAA GAGAATGCGT TCCTAGGGGG AGGICTCTAA AATGGCCACT CTGGGACTGT CTGTCTTATA TGGTTGTGGA TAAGGGATGA AATAAACCCC GGTCTCCCTT AGCGCTCCCA GGCCTATTAG GACGAGGAAA TTCCCCGCCTA GTAAATTTTA GTCAGACTGG TTGTCTGTTC TCAAACCCTG TCTCCTGATA AGATGTTATC GATGACAATG CATGCCTGAA ACCTCATTAG CAACTTTAAT TTCGCCCCGT GCTCTGCCAT TTGCCTTGTG ATATTTTATT GCCTTGTGAA GTATGTGACCA CAACCTATTC GTACANTTCC TCCCCTT

SEO ID NO:1927: (Length of Sequence = 231 Nucleotides)

CTTTTATTGG GGGCGGATAC CGCAAGGGCC CGCCCACGGT CAGGTTAGTG TTCTGCTCTT GCAGAGGCGC KACAGCCTGA CACCTCCACC TGCCACCGGC CCGGGGTTAG TGGAACATGC AAAGCTCAGA GGGTGGAGGC AGGGGTGGTC GCTGCTGAGA CCAGGGCTGN GTGCAACAGG AGGGTCAGCA CAGAGCCTGG CTGGTGTCCC TGGGCCCAAA GGGGGCTGGG G

SEO ID NO:1928: (Length of Sequence = 283 Nucleotides)

CCCCTTGCTT CCCCTGAGCC CAGGTATGTA ATTCCTACAC ACACTGATCG AGCTTGTNTG TGTGTGTATA TGTGTGTGTG
TGTGTGTNTT AATGTGACAT GCATGTACTG ATCCNGAGAA GCCTTTATAC CAAGAATAGA GCTGGGATCT CAAGCCCACC
CTCCCAAGAT CAGACAGCAG AGTGAACCAG GAGGCCACGA CAGGCCTTGT GTCARATGGC AGACGNTGCA GCAGGAAGCA
GAACCACGGG ACGGGGRNCA TGGGATGCTA TKGGCAGCCA GCT

SEO ID NO:1929: (Length of Sequence = 287 Nucleotides)

CTAGGAAGTA GGGAGAGAT TTACTAAGTA AGGAGAGAAA GGAAAAAGGA CAAACATGGA ATATGNTCAA GCAAATAACT
TCCAACAGAA ACAAGANGAT ATGITTTAAA ATATATTTCC CCTGCCCAAT AGTAAAACTT ATTTCAGGCA CAATGCATTA
CTGAGGTGAA ATTAAAGTTA CATAAAATTG AAAACATCAC ACTGGANAAC ATTTCATGGG GCTCAACTGA AGGTGGCATA
GTCCAGGAAG GCATTTGGAC ATGIATGGGG TGITTTCTTG TTGCCCC

SEO ID NO:1930: (Length of Sequence = 357 Nucleotides)

ATGGAACACT ACTGCAACAG CTCCACAGAC CGGCGGGTTC TGCTCATGTT CCTGGACATC TGTTCAGAGC TGAATRAGCT CTCTCAGAGCAC TTTTGAGGGCTG TGCACTCTGG CACCCCAGTC ACCAACAATC TCCTGTAAGA (ATGCTGAAGCA) CTCAGAGCAA AATACCCTCA TGATGTGGTG AACCACCTCA GCTGTKACGA GGCCCGGAAC

CACTACGGG GCGTGGTCAG CCTCATCCCC CTCATCCTAG ACTTAATGAA AGAATGGWTC GCCCACTTCG AGAAGTTGCC GCGCAAGGTG CTGCAGGGCA CGCGGGCTGC CTGCACT

SEO ID NO:1931: (Length of Sequence = 343 Mucleotides)

ATCACTICCC CACCCCACAG GATCIGCCCC AGAGAAAGIC CICGCTCGIC ACCAGCAAGC TIGCGGGIGG CCAAGITGAA
TGATGCIGCC CGGGGCTCTG CCAGATCCTG AGACGCTTCC CCTCCCTGCC CCACCCGGGT CCTGTGCTGG NICCTGCCCC
TTCCTGCTTT TGCAGCCAGG GGTCAGGAGG TGGCTCGGGT GTGGGCTGGA GAGGCAGAAG CCCTTTCCTG TIGGTGTCCC
AGCACATGGA GCCCCTTGGG CTGAGCACCA AGACCTTGAA CCTTTTTTGT TTTACCTTTT TTCCAAATAA CAGTTTGGAG
AAATATCAAT GAAATCTGGG GGT

SEO ID NO:1932: (Length of Sequence = 314 Nucleotides)

TTTCATGGGT TITTGITTIG TITATITCGA ATACTGAAAA AGTCCTTTGG GCTCTGTGGG GTTCCCCACG CTCACGGCTC CTTTCTCCCA CACTCACTGC CCTTCTTCCC ACAGCAAATC TATTTCAAGG ACAGTACTIT TTAAAATGAT TAATGITGAG TTCTCAACTA GCTCTGCAGA ACTAGAGGAG CTGTTTGCAT CTGTCTGTGC GGATGGAGTT TCTTTTATCT GACACCAGGT CTCCAACCAC ACTGATGCAA GGCATTTTAT CTACAGAGCT CAACTAGAAC CCCTTTTTCA TTAGGCTACT CCAA

SEQ ID NO:1933: (Length of Sequence = 378 Nucleotides)

AGCITCCIGC GGGACCACAG CTATGIGACT GAAGCIGACA TCATCICTAC CGITGAGITC AACCACACGG GAGAGCIGCT
GGCCACAGGT GACAAGGGCG GCCGGGICGT CATCITCCAG CGGGAACCAG AGAGTAAAAA TGCGCCCCAC AGCCAGGGCG
AATACGACGT GTACAGCACT TTCCAGAGCC ACGAGCCGGA GTITGACTAT CTCAAGAGCC TGGAGATAGA GGAGAAGATC
AACAAGATCA AGTGGCTCCC ACAGCAGAAC GCCGCCCACT CACTCCTGIT CCACCAACGA TAAAACTATC AAATTATGGA
AGATTACCGA ACGAGATAAA AGGCCCGAAG GATACAACCT GAAGGATGAA GAGGGGAA

SEO ID NO:1934: (Length of Sequence = 239 Nucleotides)

ATTITAAATTG ACAGCCTTCC ATTITTCGAG AAAGTACAAA CAGAACTGCT TTAGCACCCA TCGAGCCCCA AACGGGTAAG GTAAGCCAAG GTTTTAATGA CCAGCCCAGT ATCTAAGCTT CCAAACGGAT GCCAGCCCAT CACATACTYA CCCTGGGAGG CTGCTGCACG GGCATTCTCC YGATGCTCAC GGCACTTGGK GTAGGTTTCA RGATCGCCTC TTTGAGGAAG GACTTCAGG

SEO ID NO:1935: (Length of Sequence = 319 Nucleotides)

TTAATTTTT TNTCCCATAG AGGAATAGCA TTACAGTCTA ACAATCAGAA TTCTGTTACA CACATACACA GGCATGCCAC ATGACCCAGT TGAGGTGGTT GINTCCTTGA GTCTGTTGAC ACGTCACATG GTCAAAGTCT CCTCATTTCA GCCAGTCTCA ACACAAAACA CCCAACAGGG ATGCACTCAA CTTGTTGGTT CCATGTGGAA CTAGGTGGCA GGGCGAGAGG GAAAGTAGTA GAAGGGGGCT ATGGTGTGTC TGCATTCAGT CCCCTCACAT AAAGCCACAT GGATCTAGGG GGGGTATCCA AGAGCTCTG

SEO ID NO:1936: (Length of Sequence = 415 Nucleotides)

CTATTITIAC AAATTATACC TAATGAGTAA AATTAGTGTA AAGTGATAAC ATGCTTCTAC CTGTATTTCT AGTGACCCTT
TAGCGGCAGG TATTTATACC TGGTATTTAT GATGCAGTAT ATAAGTGGTG AACAATAACT GACAGTATTG TGCTTGCTGT
ACATGTCTGG TCTTTTGAAA CAGATTTTAG TAAGCATTTT CCAGAGGTAA AACTGTGTCC TTATTCTAAT TTTATTCCTA
AGGCAAAGTA GACAGGGATT ATTTCCTTGA ATCTATTTCC AAATTAATAT TTTTTTCTTT GGTATTTCTA CACTTTAAGG
CCATTTGGTC CAATTTAGAA AGTGTTGGCC TCCCTWCCCC TAGCCACATT CAAAATTAAC TTYCAAAACC TCAGGAACAG
TACAAGGAAT TTGAA

SEO ID NO:1937: (Length of Sequence = 393 Nucleotides)

TCACTCTTGT CACCCAGGCT AGAATGCAAT GGCACAATCT CGGCTCACTG CAACCTCCGC CTCCCAGGTT CAAGTGATTC
TCCTGTCTCA GCCGCCCAAG TAGCTGGGAT TACAAGCACT TACCATCACG CCCAGCTAAT TTTTGTATTT TTAGTAGAGA
TGGGGTTTCA CCATGTTGGC CAGGCTAGTC TCAAACTCCT GACCAGCGGT GATCCACTCA CCTCGGCCTC CCAAAGTGCT
GGAATTACAG GCGTGAGCAC CGCGCCCAGC CTGTNTTTCA TGTTAGATCA TAATATGATC TCACCAGATC CTTACTGAAA
ATGTACCTTA TTACAAGTAG CTAAATTTCC ACATAGAGGG NTAAAAAGAT TGGGGAATCA GGTTATGACT TTT

SEO ID NO:1938: (Length of Sequence = 407 Nucleotides)

GECCTCCTE TOGGTECAA TECAGTEGET CAGATCATAG CTCACTGCAG TCTOGAACTC CTGAGCTCAG GCAGTCTACC
TACCTCANCC TCCCAAAGTG CTGGGATTAC AGGCGTGAGC ACCGCGCCCA GCCAGAACAT CTGTTTTTAC ACCCAGAGAG
CGCCCCTCGT TAGGACAGAA CCACGGTGCC CAGAGCCAGG AAGCCGCCCT CCTGGCGCCC AGCATCTGAG CTTCTACACG
TGATGGGCGG GCTCAGGAGA GGACAGGGAG TOGTGGTGGA AGTTCCACAG CTGGCCGCGT GGGGGGCCC TTGCACCGCA
CTTGCCGCCT CCTGACTGCC CCGATCCCCG CAGCCCCTGT GCCGGATTGC ATTTYCCTCC TNTCTYCCAG GGTACTGGCC
CCAGCAA

SEO ID NO:1939: (Length of Sequence = 412 Nucleotides)

GACATGCCAC CACACCAGIT AATTITITGI ATTITCAGIA GAGATGGGGT CICACGATGC TGTCCTGGGT GGTCTTGAAC
TCCTGAGCTC AGGTGATCCA CACTTCGGCC TACCAAAGTG CTGGGATTAC AGGCGTGAGC ACCGCGCCCG GCTAAAAGAA
AGGAGATTCT AATGCATGCT ACAACACCGA TGAACCTTGA GGACATGACG TTACGTGAAA TAAGCCAGGA ACAAAAGACG
AAGGCTATAT GAATCCACTC ATATGAAGTA CCTCGATTAG CCAAATCCAT ACAGAAAGTA GAACAGTGGT TGCCGCGGGG
AGGGGGAAAT GGAAAGCCTA TATTTAATGA GTCCAGAAGC TTTTTTTTGG TTTTGTTTTT TAGACGGAGT CTCGCTCCTG
TTGCCCAGGC TT

SEO ID NO:1940: (Length of Sequence = 421 Nucleotides)

ATCCATCCC TIGCCCAGGG CCTCACATGC CCGGCTCCCC CAACCGGTCC TTCCCCTTGG GCTGCCGGTG CAGCTGTGGG
CCCAGGCTTT GGCAGGCCCA GCTTCAAGAC AGTGGGACAC AGAAAACACT TTGCAGCATC GCCTCTCCCT CCGCCACACC
CAGGTCAGCA GAGATGGGCC CCCCACCGAG AGATCACAGC TCTGGTACAG GGAGGTGGGC AGGGTTGGAG AGGAATGGAG
AGACATGTCA CCTCTATAGA AACGCGTCCA AAGTACAAGC TAAGCAGGGG GAAGGAGGAG GGCCAGAGAG CAGCCGGAAA
GAAGAAAAGA GGAACACGGC AGGGGTTCT KGGGGAGGAG GGCCTCACAM CACCCCGCAG ATGAGCGTCT TCACCACGAA
GGTGTTCTTC GAAGTKGCGG T

SEO ID NO:1941: (Length of Sequence = 377 Nucleotides)

GTCAGCTCTA GAGGCACCCT GCATCATGCC CACCAGGGTG ATCCCCCTGG GATNGACCAT CTCGGGATAT GAGGCCTCGG
AGGCTGGGGT TGAGATTTGG TCCTGAAGAG CTTATAGCCA GATTGCCACA TTCAAGTGTA AGTCCAGGAA AGGGGCAGGC
GGCAGTGCAC AGGGATTTAT CAGTTCCAGA ACCTCACAGT GATAAGAGGC TTTAGAGAGC ATCTAATCGA GACCTTTAAT
TTTTCGGGGA GAGCAGCTGA GGCCGTGTGG AAAATTAGTG GAGAGCTGAC AAGTGTCTTG GCTCCTGGCC CAGGGGTCCG
TGGTCCANCA CGTTGTCGTT CAGTTGGAAG CAAAGGGCTT GCCCGTGATT ACCTTCC

SEO ID NO:1942: (Length of Sequence = 401 Nucleations)

TGAGAACATT AAGAAGGACA ACAAAATTAA ACATTCTTTA ATAAAATTC TATAGAAAGC TCAGTCATAG GCGAAATACT
CATTTCTCTT TCCCATATCA CCGAGGATTG AGAGCTCCCA ATATTCTTTG GAGAATAAGC AGTAGTTTTG CTGGATGTTG
CCAGGACTCA GAGAGATCAC CCATTTACAC ATTCAAACCA GTAGTTCCTA TTGCACATAT TAACATTACT TGCCCCTAGC

ACCCTAAATA TATGGTACCT CAACAAATAA CTTAAAGATT TCCGTGTGGC GTGAACCATT TCAATTTGAA CTAATATCCT TGAAAAAAAAT CACATTATTA CAAGTTTTAA TAAATACAGT AGAGAGCTGG CATTTTTCTA AATACTGGAT TTCAGATCTG G

SEO ID NO:1943: (Length of Sequence = 351 Nucleotides)

CAACTCAGGT TAGCAACTGC AGGAAAACTT TCTTCATTIT CACIGAATTI TAAAGAGAGA ATCCIGICIC TATITCTCAG
AGAAACTTAG GTGAAAAGTA AAAGAGAGGC AAAATCTCIT TCCITCATGA GATACITITA TITTITATCIC TITCICTACT
CATGTGCTTA ACTGGIGAAA TGATTCTGTA GAAATAGATC CITCIGATTC TGCATCTCAT TTCCITATGG CAACTACAAC
AGGAGGAATC CAGCTGGAAA TGCCACTAAC CCCACATCCA GCACCTGAGA GAGGAAGCCA GTCGGAGCGC CGTGCTGGGC
TCACTCACTC TGGCCTGCGC ACTGGGGTTG T

SEO ID NO:1944: (Length of Sequence = 406 Nucleotides)

GCCCAGGCTG TCTCAGAATC TTGATGGGGT GGTCATTGAG CTCCTCTTCC GCCAGAGCAA GATCAGTGAA GTCCTGGGAG GCAGTGGCTA CAACTCGGAC CGCCTGCCC TGCCCTACAT TCCTCAGCTG ACAGATGAGG ATCGTTTATC CAAGAGGAGG AGCATTGGAG AGAACATCTT CCCTGAGGAT CCCGAGGATG GTCTGGTGAA GACCAACATG GAGAAGCTGA CCTTCTATGC CCTCTTAGCT TCAGAAAAAAC TTGATCGTAT TGGCGCCTAC CTCTTTGAGA GGCTCATCCG TGACGTGGGT CGNCATCGAT ATGGGTACGT GTGCATTGCT GCTCATGGCC TGCCACTGCC AGAGCATCAA CCTCTTCGTG GAGAGCC

SEO ID NO:1945: (Length of Sequence = 362 Nucleotides)

TCAAATTGTG AAATTNAGAA TTCTGCTATG ACAAGTGGAA AATTGAGAAA AGACGCAGAG CCACTTTTTG TNATCGTGTA
GGTGACAAGG AGTCTCCCAA GTATATCCTG CTAATAGGAG TAGCTCTCAA AAGTTAATCT CAATAAAGCC TCCTAAAGTC
TCTGGCCAAAG AAAACTGCTG CAATCCCTTG TGCAATTCTC CAGACTAAGC TGTATGGGGG AAGCCTACCT TTTTTCAGCC
CGAAGTTCAG GAGACTGAGG ATGTAACTGG GGACATGATC ATTGNTTCAA AGGTGATTGC TTAAGTATCT TAAAAATGTA
TAGAGCTAAT CTGAGTACCG CTTAAATTCA AGAGCCGTGG CT

SEO ID NO:1946: (Length of Sequence = 408 Nucleotides)

AACCICINAC CCCCAGGITC AAGCAGICCI CCCACCICAG CCICCCGGGI AACIGITCIT TGIAACICIC TCATCATCGA
GGCIATATAT TAATAGACAT GGIATTAAGC CCACACGAAA CATTCAGAAT TAGAATTGGA TTAAGAAGAC GCGITTTGGC
ATCACGCIGA CTACTCCICA TCTCCGICCI CGGGGGGGI GATGCCAGGG TGGGACTCIT TGGAAGGGCCI ATCAATCACA
GGIGCGCTAA AATCAAAAGG TGGGICAGTA GGITAGGGAG GCACGCGCGA AAGGAGATGC CAGCGGGIGI TAAGAAGGAT
ATGGICAGAA GAGCTCTITG TCTCCATCCA CGGGGCCTCI GCTCAGCCCG TGITGICTCG GTGAGTAATT CGGGAGCAGT
GCACGGCT

SEO ID NO:1947: (Length of Sequence = 426 Nucleotides)

CCATTIGACA CIGITACIAT CIGCAACAGI TCITGCAGTA GAGGATGCAC TICAAAGTGC ACTGCITTAC TGICTCACTG
GAATTCIAAA AATCIAAGCI ȚITATCITITI AACATTAAGC TGIGIGGGAA TGIAGCAACC TCCTGGGTGG TGGGGTGGGG
GGCATCITCA ATTATITAGG TCICACTGGA AAGITTGAGA TCAGAGTITIG GTAGGTGGTG TAAGGGGACA ATGAGTAAGG
GAGAGAAAAT ACAGGACTGA CITGGGGCAA AAAACGCCTG ATAATAATITI GTGAAGCACA TITTCAAACT CATTTATTCC
TTACAAGGAT CCTAAGAGGC GGGTATTATG TCCNGGTTAT ACCTGGAGGC TTAAATTGAA GGAACATCIN CAAGGGCACA
CAGITTAATG AATGGCTGAG GTAGGA

SEO ID NO:1948: (Length of Sequence = 349 Nucleotides)

SEO ID NO:1949: (Length of Sequence = 378 Nucleotides)

TTCATTCCTG ATTITATCCC AGCTGTCGGG GATATTGATG CATTCTTAAA GGTCCCACGT CCTGATGGAA AGCCTGACAA
CCTTGGCCTA TTGGTATTGG ATGAACCTTC TACAAAGCAG TCAGACCCTA CGGTGCTCTC ACTCTGGTTA ACAGAGAATT
CTAAGCAGCA CAACATCACA CAACATATGA AAGTAAAAAAG CCTAGAAGAT GCAGAAAAAGA ATCCCAAAGC CATTGACACG
TGGATTGAGA GCATCTCTGA ATTACACCGT TCTAAGCCCC CTGCGACTGT GCACTACACC AGGGCCATGC CCGACATTGA
CACGCTGATG CAGGAATGGT NCCCGGAGTT TGAAGAGCTT TTGGGCAAGG TAAGCCTG

SEO ID NO:1950: (Length of Sequence = 357 Nucleotides)

TCACTAACIT TACGAATGAA AGAAAACAAT TCCATCCCTC TCACAAAAAG GACATCITTT AAGCTTTCCT CCCAATCIAA CCTCCATGGG ATCTCAGAAA TTCCAATTCT TATAACTCAA ATCCCCACAG TGGTGTAGAT GCATTAACTC CCCGGGGACA GCAATCIGAG GCAGGCAGGT TCATTAAACA AACATGTTCT GTGCCCTCTG GCAGAGAGGG CAGCAGGACA TGCACTGCCC CTGAGCCAAG CTGTGGCATG GGCAAGGACA TCAAGTAGCT GACAACGGTC TGTCCATCTC AGCTGGGGCA GAGGGGCCAG TTCAGCCTTG AAACAGCAGT TNGGGAGTGT CTCAGCT

SEO ID NO:1951: (Length of Sequence = 336 Nucleotides)

CTATCTCCCC AAATCTACGI TTCACCATTI GIACIGITAT TITTITAGCC CAAGCCACCI TTATGTCACT CCIGGAACAT
AATAACTGCI TTCTCACTCA TCTCCTACAT TTINACCICI TATAATACAG TCCACCTIGI ACCGAGCAAC AAGAGTTATC
TTTCTGAAAT GCATATTAGA TCATGTCACA TCTCTACTTG AAGCTCTCTA AAGATTTCTC ACTAAAAGCG AAGTCTAAAA
TTTCCACCCA GACCTATAAG GNCCTTAAAT GATCTTACCT CTCTACCTAC CTCTNCGATC TTACCTATCT TCAACCTCGG
TTCTATTTTC TATATC

SEO ID NO:1952: (Length of Sequence = 413 Nucleotides)

CAGTATGIAA TITAATCAGC AAATGCCCCA TITCCATCTC TACCGGAAAG CITTCAGACG CATTCCCAGA TCAGACAGAG
GACTAGGGIT AAGGCIGGGA ATGAAACACC AGCTAGTATC CCAGTGAGCI TICCCAAACA CACATACACA GCAAGTCAGA
CIAAACAACG TCCAACIGAA GACTCACCTC AAATACITAG ACCTAAGATT CACGTCCAGG CTCTTTCAGA TACACCAGGT
AAGTAAGCAC TIGGCATTCC TATCTCAGCC ATTCACTTCA CAGAATCITT TGGGIGCCTA CIGIGIGCCC AATACTGTGC
TTAGGGGTAC TIGGCCTCAG CAGGAAAAAA AATTAAAAGT GITAAATGIT ATGAAGGAAC AGATTGGNAT AGGAATCACA
AGGCATTCAG GTC

SEO ID NO:1953: (Length of Sequence = 382 Nucleotides)

GITTCACTCT TGITGCCCAG GCTAGAATGC AGTGGCGATC TTGGCTCACT GTAACCTCTG CCTCCCGGGT TCAAGTGATT CTCCTGCCTC AGCUTCCCTA GTAGCTGGGA CTATAGGTGC ATGCTGCCAC ACCCAGCTAA TTTTTTTTGTA TTTTTTAGTAG AGACACACTT TCGACATATT GCCCAGGCTG GTCTTCAACT CCTGATCTCA AFTGATCTGC GTACCATCA TGTTTAGGGC GCTGGGGATTG CTGGCCTGAG CCACCGCACC CTGCCTAGAA CATGCTTTIN AATAGTGTCT CTAACCATCA TGTTTAGGGC

CTTAGTGCTT ACCTCTTAAA GAAGGGCTGC TGTTGAGGAT TCCNTGAGAT AGTGTTTGAA AA

SEO ID NO:1954: (Length of Sequence = 389 Nucleotides)

GGAAAAGCGG GACCCAAACA GTGGTGCTGG GGAAATTTGT TCCTGTTCCC TTTGGAAGGC TGAGTGGGTG ATGCAGCACA
GGAACAAGGC TTGGACGTCA GAGGTCTCAT CTTCACTGTG ACAAAGCATA AAGGACTTGG GGTTGAGCGT GTGINIGGGC
TCAAGTGACC ATGCAAGINC TGTCACCTCC TTCCTAAGAC CCCATCCTTC TCCCAAGTCC TCCACAAGAG CTACCTTCTT
CAAAACAATA ACAGAAACAC ATCAAGNTIN GCGTCACTGA AATTGAAGTT CTGAATTCTG CCGTCACCCC AGCAACAGTG
CCAGTTATGA TGAGACACTT GACCCAGCAC TTGGGTTGAT GTCTTTGGCT GTTACCGTGG CACCTAGGT

SEQ ID NO:1955: (Length of Sequence = 277 Nucleotides)

GCCTCIAACT CCACGGCTCA AGTAATCCTC CTGCCTCAGC CTCCTAAGTA GCTAGGACTA CAGGTGCACA CCACCACACC
CAGCTAATTT TITINCITTT TGATTTTTGG TAGAGATAAG GTCCTACTAT GITGCCCAGG CTGGTCTGAA ACTCCTGGCC
TCAAGTGATC TGTCTTAGCC TTCTGAGTAG CTAGAACTAG TTTTAATGAC CNAAAGAATT ATGTGTTCAC CNGTGATTTT
ATGTGTTTTG TTAAGACATT CAGAATTTAG AGAAATG

SEO ID NO:1956: (Length of Sequence = 380 Nucleotides)

GTGTAATGIT CTGAGGGTGG CGAATGCAGG GGCGCGTTCC TCCCGCTGTC GATCTGGAAC ATCITCTCGC CAACAAAGAG
CAGGGTGAAG ATGAGGGCAA GCTGGTAGAC AGCATGGCCC AGGATGTTCT TCATCATGGT CCTGGAGATG AGCGGCTTGT
TGCGGCCGTA CGGTTTCCTC AGCAGCAGGG TCTCCGTGGG CGGCTCAGTG GCCAGTGCCA GCNAGGCAAA CGTGTCCATG
ATGAGGTTCA CCCAGAGCAT CTGCACGGCC TTCAGAGGGG AGTCCTGCGT GATGCAGGCG CCTNTAAAGC CACAATCACG
GCCACCACGT TGACGGTGAA GCTGGAACTT CAAGAATTTN GAGATGCTGT CATAGACGTT

SEO ID NO:1957: (Length of Sequence = 328 Nucleotides)

TGIGATGITT CITITITAGC CIGITGATGI GGIGAATTGI ACTGATTGAT ATTTGAATAT TAAACTGGCT TTGCATCCCT
AGAATATACC TCACCAGGIC ACTGIGTACT AGGITGGIGC AAAAGIGCIT GCCATTTIGG ACCATGAATT TTGAATCATT
AAAACTAGGC TCAAACACAT CIGITATTAAT CAAAGIAAGA ACCATTACAA TCAACACAAT TTTGCCAACA AGAAATAAGT
TTGTTTACTC CIGTAGCATA AAAATCCGTG CITTGAGATT CGAGGAACTT TTGGAAAGCA CITTCTGCAT CCTGCTGGTT
GTGGAAGC

SEO ID NO:1958: (Length of Sequence = 254 Nucleotides)

CTAGAAAGTA TCTTCTCTTT ATTTAAGTTA AACAATTTTC AAGGATGGTT TCCATCTATA AAATGGACAA AGTACAAGCT CTGTACAGCA GTTCTTTTTA AAAATCAACT GGAAAAAAAA ATTACCAAAC TATATTTTGA ATTTGCAAAA CATACTCACA GATACCATCA TCTGAGCTTT TATGAGGACA TAAGAAAGGA CCACCACAGA GAAGACAACT AACTTCGGCA CGCTTTGCTC GAAGGGGCTCT TAGG

SEQ ID NO:1959: (Length of Sequence = 259 Nucleotides)

GTAATACGAG AAAAATCACA ACAGAGTAAT AAAGATATAA AACTITCACA ATTAACACTC ATCAGTGTGA TAAACTAAGC CCATGTAAAA GTAAAAATCT CTCACAGTTA ACAAACGTCT TTACTITCAC TAAGAAGGAA CTGAAATTAA AGTCCTTAGT CACTTTGGAG GTGGCTGCAA AAGCTCACAA CATAGTTGAT CCTTAAAATA ATTATGAATG GCAACCAGTG CTGCCTTTCT CTACTCAACC ATGCAACTG

SEO ID NO:1960: (Length of Sequence = 329 Nucleotides)

GACTACAGET GIGCCCCACG ATGCCIGGCT AATTITTAAG GITTITGIAG AGATGGGGIC TICCIATCIT GCACAGACIG
GIGIGGAATT CCIAGCICAA GCAATTITCC IGICICAGCC TCACAAAGIG CIGGIATTAC CCGIGIGAGC CACCGIGCIC
AGCCCAGICA IGIATITCIA ATTATIGIAT ITGIGAACIA ATCIATGAAC AACAAAAACA AACAAACAAA CAAAAAGGGI
GGCATITCIG GGCCACCAGG GAAGGIGGGA TIGGGGIIGC AGCIATITIC AAATTATATT AAAAGCAGGA TCCCAGITAG
AGCGCIATC

SEO ID NO:1961: (Length of Sequence = 282 Nucleotides)

ATCCICCAC CICAGCITCC CAAAGIGGIG AGATTACAGG NICGAGCCAT CGCACCCGGC CCAATTATIC TITCIAAACC ATTICCICIT CIGIGITCAT GCCITTAAAA ATAAAATTAA AAAAAAAAA AAAAAAAATC CITAAAATTI CICAGGIGIT TICCATATCA TITTATTATC AAGAATAIGG CIAATCAGAA GICACAGCCA GCCCCGAAC TACAACTACA AAACAIGCAT ATTATAGGCT ACACTGAGGG ATTICIGAGG TIAGCAGATG CA

SEO ID NO:1962: (Length of Sequence = 328 Nucleotides)

TECTGGTGTC CCTGCTGTCA TCCTCAGGAG GCCAAATCAG TCCCAGCCTC TCCCACCATC TTCCCTGCAG CGATTTCTTC
GAGCTCGAAA CATCTCTGGC GTTGTTCTGG CTGACCACTC TGGTGCCTTC CATAACAAAT ATTACCAGAG TATTTACGAC
ACTGCTGAGA ACATTAATGT GAGCTATCCC GAATGGCTGA GCCCTGAAGA GGACCTGAAC TTTGTAACAG ACACTGCCAA
GGCCCTGGCA GATGTGGCCA CGGTGCTGGG ACGTGCTCTG TATGAGCTTG CAGGAGGAAC CAACTTCAGC GACACAGTTC
AGGCTGAT

SEO ID NO:1963: (Length of Sequence = 277 Nucleotides)

CCAAGAGACA CCCCCCGCAC TCCTGTGCCC GAGCTGTCTC ATCTGTGATT CACAGTCTGC TCTTTCTGCC TGCTTGTCGT
GAGAAGTGAT TTTNAACCCC GAGGTTAGAA AGGGAGCTAT TTTTGAGCTG CTTTTTTGTTA AAAGGCAAAT TTTCTGCTGG
GGACTGGCTT TACCCCGTCT ACCTAAATCA TTTCTTACTG CCTCCTGTAA CAGTCGCCTT TTGTGTTCTG CTGCNATTTG
TTTGAACACA GTCCACAGGT TCAGTGGTTN CATCTCT

SEO ID NO:1964: (Length of Sequence = 230 Nucleotides)

CAATGCAACC TTITAATTCC AAGCAGAGTC CCCCTCCCCC AGCATGGTCA CACACAGT GGAAAGGGAT GTCAGGGTCT GGGCAGGAGC AATACCCAGA CCTGGGCAAA AATATAGATA TCATTATATA CACACGTGGA CTGGAAAGAA GTCAAGCTGG GGGTGTAAGG TAGGGCAGGG GCAGGTGAGG AAAGCAGCTG GGGGGCCCCC AATAAATTAC ATTCTTGAGA

SEO ID NO:1965: (Length of Sequence = 299 Nucleotides)

CGCCGTGGAT CCCGAGAAGG CACAGCAGAT GCGCTTCCAG GTGCATACCC ACCTTCAAGT GATTGAGGAG AGGGTGAATC
AGAGCCTGGG CCTGCTTGAC CAGAACCCCC ACCTGGCTÇA GGAGCTGCGG CCCCAAATCC AGGAACTCCT CCACTCTGAA
CACCTGGGTC CCAGTGAATT GGAAGCCCCT GCCCTGGGG GCAGCAGCGA GGACAAGGGT GGGCTGCAGC CTCCAGATTC
CAAGGATGCA GACACCCCCA TGACCCTTCC AAAAGGGTCC ACAGAACAAG ATGCTNCAT

SEO ID NO:1966: (Length of Sequence = 320 Nucleotides)

GTCCCTGCAC ATGCGTCTGG CAAGACGGGT CAGCTTTGTG GTCTGAAGCA GGAAAGTTTG TCTGTNCTTA GCCAGTAGCT
TGGCCCTGTT GGCGCTGGTT GTGTAAGGAG ÄGAGACTTTG AGCTTCAGGT CTGGATAAAT NACCCCTTGA GTGTGGCTCC
GTGGTGCCCC GAGTGGCCCC CTCTAGCTGA GTTGGGGTCT TCAGTCCCCC ATACTTCTC CAGTAGATCC AACTAGACCA
ACAGAGGCGG CACTGCATGT TAGGTGGGCC CCAGGCATAC CACTGAGCAG ACTGTGTGGT GTGGCAACTC TCACAAGTCA

424

SEO ID NO:1967: (Length of Sequence = 296 Nucleotides)

GCTCTGCTGG CCGTGCAGAA GCTCATGGTG CACAACTGGG AATACCTTGG CAAGCAGCTC CAGTCCGAGC AGCCCCAGAC
CGCTGCCGCC CGAAGCTAAG CCTGCCTCTG GCCTTCCCCT CCGCCTCAAT GCAGAACCAG TAGTGGGAGC ACTGTGTTTA
GAGTTAAGAG TGAACACTGT TTGATTTTAC TTGGAATTTC CTCTGTTATA TAGCTTTTCC CAATGCTAAT TTCCAAACAA
CAACAACAAA ATAACATGTT TGCCTGTTAA GTTGTATAAA AGTAGGTGAT TCTGTA

SEO ID NO:1968: (Length of Sequence = 311 Nucleotides)

ACCCCCTICA CICCCICCA CCAGCICIGC AGCCAGCCIA TGGCAATTAT ATTITAAGAG GIGITCCCAG GACITITGGG
ACCTACIAAA ACAATGATGG TIATTITAGA TGTGATGATT TATATTITATG TAGAGATATT TCIGGACCAC TCAAGCTCIT
CGATACCAAA ATCAGGAGCA TCITGGGATT TATTAAATTA TGTAAGAAGA TAGCACAGAT ATCGGGATAT TATTGTGTGA
AAATGCTGCT TTTACTITGA TGTGATCTCA TTGATGTACA CAACCAAGTT CCAATAAAGT GCTAGAATGT G

SEO ID NO:1969: (Length of Sequence = 266 Nucleotides)

CAATAATAAA AAGGATTATA TTCCTGATAC ATGCAATATG GGTGAACCGT AAAAATATCA TGCTGAGCAA GAGAAGCCAA ACACAAGAGA ACATGTTGTT ATGATTTCAC GTACATGAAA CTTTAGTAAA GACAAGTCTA ATCCATAGTG ACAGAAAGCA AATCAGTAAC TGCTGACAGG GGCAAATGAG GNGATGATCT CAAGGGNACC TTCTGGGGTA AGACGCTGTT CTGTATCTCG ATCCNATTGG TGGTCACACA AGTGAA

SEQ ID NO:1970: (Length of Sequence = 317 Nucleotides)

CTCGGGAGGC TGAGGCAGAA GAATGGTTG AGGCCAGGAG GCGGAGGTTG CAGTGAGCCA AGATTGCGCC ATTGTACTCC
AGCCTGGGCC ACAAGATTGA AACTTCATCT CGGGGAAAAA AAAAATGAGC TAAATACAAG AGATGGTAAT GCAGGAAATG
AGAGAGAAAG AAGCTATAGA ATGCACCATC AGTCTTTGCT GAGAGGAGAA GCTAGGACAC TTATGCGCAT GTNCCTGTCT
GCCTTCCTTC CCGTCCCGCG GATGGTTGGA GCAGGTCTTT GTTTGCTGCA GAGCATGCCA TGTCATCCTC CTTGTCT

SEQ ID NO:1971: (Length of Sequence = 263 Nucleotides)

GIGCATACTG CTGAGGCGGC TACGCTGGCA GGGTAAGCAA AAGAAGCACC CCAGCCTAAG TTTACAGAGA ACCAGGACAT CATTTTGAAT ATAACTTAGT TCTAATAGTC AAATGGCCAC TCAAGGTGAC AAATAGGAAC TTCAGTGGTC ACCCCTCGGA AGCAAGCTTT CAATGTCCCC CACCTGTAGA AGGCTGAAAA ACATCCTCCA AAGATAACAG GTTCCAATCA CTGGAACCTG TATTACTTAT TACCATTAAA TAT

SEO ID NO:1972: (Length of Sequence = 295 Mucleotides)

GACAAAGAA GCAGAATAAT TITACCIGAG AAGAAACCAG GAGGCITCIT CITCTTCTTC TCTCTCTTT TITTITTTTT
TITTIGACIA TACAGAAGAA AACTATCAGA GITAGGITAG AGAGTIGGGT TIGGGGTCAG GITGIAGCAT GIGITATATT
ATGGGTTAAA TIGGGTCCTC CCCAAAATTA ATAGGTCAA GTCTTAACTC CCTGTACCTC AGAATGIGAC CNCATGGGGA
AATAAGGTCA TIGCAATATA ATTAGGTAAA ATAAGGTCAT ACTAGAAGAG GGTAG

SEO ID NO:1973: (Length of Sequence = 243 Nucleotides)

425

SEO ID NO:1974: (Length of Sequence = 304 Nucleotides)

GGATGAGATG ATCGACGTCA TCGGGGTGAC CAAGGGCAAA GGCTACAAAG GGGTCACCAG TCGTTGGCAC ACCAAGAAGC TGCCCCGCAA GACCCACCGA GGCCTGCGCA AGGTGGCCTG TATTGGGGCA TGGCATCCTG CTCGTGTAGC CTTCTCTGTG GCACGCGCTG GGCAGAAAGG CTACCATCAC CGCACTGAGA TCAACAAGAA GATTTATAAG ATTGGCCAGG GCTACCTTAT CAAGGACGCC AAGCTGATCA AGAACAATGC CTCCACTGAC TATGACCTAT CTGACAAGAG CATC

SEO ID NO:1975: (Length of Sequence = 233 Nucleotides)

CCTTCTCCAT CACCCTTGGA CCCTCTCTGA GTGGTCTCTC AAGGCACATT TATTTTCTCT GCTGCAACCT ACCAGATCTG
ACATCCACCT CCCCCAGCAC CCATGGGCCA AGGAGGCCTG GGGCAGCCAA GGGGAGTTCC AGGACCAAGC AAGCAAGAAA
CCGTTCTTTG AACACATGGT TAAGCTTCTT CCAGCATGGC CCTAATTCCC CTACCTGCCT AAGCCAGGGG AGT

SEO ID NO:1976: (Length of Sequence = 162 Nucleotides)

SEO ID NO:1977: (Length of Sequence = 270 Nucleotides)

GECTGAATTA AGAGCATCCA GAAAGCCCAG GCCCTCCATA GGCTGTGGGG GGATGATCIT CACTITGATC TCTTTGGTGG
CATTAGGTGT TGTGTTGAGT GGCTTGTATT TCTTCTCTGC AGGGGAGTG GCATCTCCTG GAGCAGCTAC GTTGCTCTGA
CGTTTGAGGG GGATGGGTTT AAGGTTGTAC TTGTCAGAAA CCACCACTGT GCTGGCATTC TTCTTCACAG GCACCAAGGA
TGGTGTCTCC AGCTCTAGTC CAGTGGAACG

SEO ID NO:1978: (Length of Sequence = 167 Nucleotides)

TTGCAGGAGT TGCTGATATT TATTCAAACG TCATCCATAC AATAAAGAAC TCTNCTTTTA AAATTCCATT TACATCAGCA GTTAAAAAAA AGTGACAGTG GATGAAACAT GANGCTGTAA AGTGCCTTTA TGGGGAATNC AGCCACGCCT GCCTCCACTG TGCTGGG

SEO ID NO:1979: (Length of Sequence = 346 Nucleotides)

CATCATAGCA ACAAAGGGCT ATGTACTATA CTCAGGAAAA CCATTTATTT GCACTGGAGG CAACTGTTCT TGAGAGAGGA
AAAGTAAATT GTCCAAGATG TAACATCTTA TAAATAGCAA AGCAAGGATG AAAATTATTA TATTTNACTA AATCAGTATG
AGAATCCTGA TTCTTCATTA TTATATCCCC AACACTCTAT CAGTTTGTTG AACAAATCAA CAAATAAGCT TGAATAAAGG
MTCCACATCT CAATTCTCCT CCACCATTCT ATATTGCCCT TCATCCCTAC ATTAAAATCAN TTATTTCTGC TTTTTTTTCTT
TAACAATTTA TCCCTAAAGT AACTAG

SEO ID NO:1980: (Length of Sequence = 174 Nucleotides)

CACAAACTGA CAGAGGAGAC AGGAGGAATT TAATATTACA TGCTATAATG ATATTTATCT CACAGTTTAT ATTTCATTCA
TTTATATTAT TTTTTTAAAA GGTTTCTTTA TCAGCTACTA AACATCTCAG CAATTTGGTG TGCATAGCTC TAGATTAAGC
AACAAAGAAT TGTA

SED ID NO:1981: (Length of Sequence = 276 Mucleotides)

TGGNICACTC ATAAGITITC AGIGGITAAT TACIACAGIT TAAGAAGACG TGIGATITAT TITTAGATCT GACCCAGCAG ATCATACCIN TNCNIIGAAT TACAIGGICI TCITITGGCI TCITAAGATGI CACACTCCIG TCITAGIGGC CACTGCICCT CAAGCCCCCT TIGCTAGCTC TICCTCATCT GICCAGCCCT AACCIGACCG TGCTATGTAA GTCTTCTCCG TNITCACCCC CINCCWGGT GACCGITATA CINCCAAACC TACAGG

SEO ID NO:1982: (Length of Sequence = 288 Nucleotides)

GCTGCAGAGA GGITGINTCC AGGAGCAGGC TITCCCGCTC GGGATCCAGG TCATCCCCCA CCAGAGAAAT TICACAGCCA TCCAGGITGT GCACAATCTC ATCCGACATG CGTGTNTCTG TCACTGTGCC CTGCCAACTC TCATCCTTTT TGGCCTCCAC CTGGTGAGAA ATGGAGCAGG TGATTTGAAG ATCAGGGAAC AAAGGGACGC CGTTGGTTCC CTCAAAGTCC ACAGCTNGGC GGGCAAAAATG AGCAGTGCCA CTCAGCAGGA TCTGGGGGGC GTCAGGCT

SEO ID NO:1983: (Length of Sequence = 273 Nucleotides)

CACAAGCCAC TITCAGCCTC CAGTGGGAAG GCTCCAGCCA CACGCCGATA TITCGTCCTG CTTCCCGTCA TCTCATATCT
AAAAGTCATG GCTTAAGTTA GGCAATAAAA CCTGTGGCTT TAGGCATCTT TAGTAAAAAA GCTGAACAAA TCCCAAATTT
ATTCCCATTT TCTTGAGAAA TAAACTTCAT AAAACAACAG ACAGCTGTCA TGATTACTGA GTTTTGGCTG ATGGCGAAAT
AATTTTTATG TAAGTATACT GAATAAACAT ACA

SEQ ID NO:1984: (Length of Sequence = 221 Nucleotides)

GAAGAGGCTG CTCTGGCCTG GGACACCCCC ACTGCTCTCA AGGAGCTGGC ATCTCAGTGG CCTCTNAGCC CAGCCTGAGC
CCTGTGGGAG TNCGGGGGCA GTGACTGGAA TGTNCTGCTG GGCAGGCCTGC AGCAGCCGAG GTGGCCCCAG GGCAGAGGAG
TGCAGCGCAN CTCATGGGTG CCCTATGCCA CCCCTGGTGC TCACTGGGCT GCTGATGCCG T

SEO ID NO:1985: (Length of Sequence = 197 Nucleotides)

TTGCTACCAT GAGGGAAGTG CTCGTTGCTT GGCCTACAGC AAGINATACA GCCTGCGAGG CACAGTCCCC AAAAGTCTAG CTGCAATTCT ATTGGTGGTT TTCCCCAAAC AGCAATAACA AGATGTTACC TGGAAGCACA CCAGAGCCAA TCATGACTCA GGCCTGTCTA GATGTTTAGA TGTCTGGAAA TATATTT

SEO ID NO:1986: (Length of Sequence = 268 Nucleotides)

<u>SEO ID NO:1987:</u> (Length of Sequence = 282 Nucleotides)

GTCCTCACTG TAAACAAATG AGGATGGAGG ACACTGAGAG GNTCAAATAT GAAAGGCAGT ATGGGGAGTT AGAGCCACTC
GTCTACTCCT GTAAAGAGCA TGACTACTCA CAGTCTTTCT AGCGGGTAGT CACTCTTTCA TTTAACAAAT ACTTAGTCCC
TGCAATGATC TAGGATAATA ACTCAACAGT GTATATCAAG AGCCTTTAAA AAGTTATACC TGGCCGGGCG CAGTGGCTCA
TGTATGTAAC CCTAGCACTT TGGGAGGCCA AGGCAGGCAG AT

SEO ID NO:1988: (Length of Sequence = 226 Nucleotides)

TABEGGGT TOGGTOTOTO AGGAAGTTAG GOCATAATTT CTGCAGGTTC AGTGATTAAC TAGGATOCAT COCATGCTGT CTTGAACTGT TCAGGAATGG GAAATTCTCT ATAATCACCA TCCTGAGGGA TAAGTATGTT CATTTCAGAT, GACTTGGCGC TCACGNTCTC ACAGTCTAAT GCATCTTCAC TGAGGTATAT GTGGCAACCT TCTGTCTTAT TAATGG

SEO ID NO:1989: (Length of Sequence = 193 Mucleotides)

CTCCCTGTAG GTCATGTCCT TGAGAGTTAA AAGATGGGTT GAGTAGGCAG AGGTCTCAGG CACCGGGGAC AGAAGACAAG GACATTCAGC ACGGGCAGCC ATGCTCTTCC CAGCACCCAG AAAAGGCCCA GGGCCCGGAC TCCTGGGTGT GGTCATGAGA AGGGCCTCCG ATTCAGCCTC TTCTCTTCTT GTG

SEO ID NO:1990: (Length of Sequence = 223 Nucleotides)

CTECTTCATT TACCACCACC AGCCGATGGA CCAGTITATT GGATTCACCT ATGATACCAG GACTITTCCA TTCAATTCAA
TTCAACAAAC TITTAGAGAT CGCCCCTATT CCAAGCTCAT CCAGGTTCTG CTTCATGAAG GCAGGCTTTG GCATATCAGA
CATAAAAAAGC TGGAGGAACT TGAGGATTCT TITGTGGGTA AGTATATAAA GTGCATTCCC ACT

SEO ID NO:1991: (Length of Sequence = 385 Nucleotides)

GCAGAGAAAG TGCCAGGCAT CAACCCCAGT TTCGTGTTCC TGCAGCTCTA CCATTCCCCC TTCTTTGGCG ACGAGTCAAA
CAAGCCAATC CTGCTGCCCA ATGAGTCACA GTCCTTTGAG CGGTCGGTGC AGCTCCTCGA CCAGATCCCA TCATACGACA
CCCACAAGAT CGCCGTCCTG TATGTTGGAG AAGGCCAGAG CAACAGCGAG CTCGCCATCC TGTCCAATGA GCATGGCTCC
TACAGGTACA CGGAGTTCCT GACGGGCCTG GGCCGGCTCA TCGAGCTGAA GGACTNCCAG CCGGACAAGG TGTACCTGGG
AGGCCTTGAC GTNTGTNGTT AGGACGGCCA GTTCAACTAC TNCTNGCACG ATGACATCAT GGAAG

SEO ID NO:1992: (Length of Sequence = 312 Nucleotides)

GGCTTACAGG ACAGAAAGGT CCCTTCTCAC AGTITGGGAG GTCCGAAGTC TGAAGTGAAG CTGTCAGCAG GGCCACACCC CCTCTGGATG CTCCAGGGGA GGGTCCTTTG CCTCTTCCAG TTCTGGTGGC TCCAGGCATT CCTTGCTTTA TGGTGGCATC ATTCATCTCT GCTCCGTCTT CACGTGGCCT TCTCTGTGTT GTCAAATCTC CTTCTCTGTT CTCTTGTAAA AACACTCGTC ATTGGGGATT AGGGCCACC CCAATCTAGA TGGTCTCATC TTGAGCCTTT ACTTTAGTTA CCTCTGCAAA GA

SEO ID NO:1993: (Length of Sequence = 429 Nucleotides)

CTGTTTTAC TCGACGAGGA GAAGACCTT TCATGTGTAT GGACATACAG CTCGTTGAAG CACTGTGTGG CTTCCAGAAG CCAATATCTA CTCTTGACAA CCGAACCATC GTCATCACCT CTCATCCAGG TCAGATTGTC AAGCATGGAG ATATCAAGTG TGTACTAAAT GAAGGCATGC CAATTTATCG TAGACCATAT GAAAAGGGTC GCCTAATCAT CGAATTTAAG GTAAACTTTC CTGAGAATGG CTTTCTCTCT CCTGATAAAC TGTCTTTNCT GGAAAAACTC CTACCCGAGA GGAAGGAAGG GAAGAGACTN ATGAGGATGA CCAAGTAGAA CTGGTGGGAC TTTNGATCCC AATCAGGAAA GACGGCGNCA CTNCAATGGG GGAAGCATAT GAGGGATGAT GGACCATCAT CCCCAGAGGT

SEO ID NO:1994: (Length of Sequence = 377 Nucleotides)

TGGGGTTGCC AAACCAGTTG CCCCTGTCCT GTGTCAGCCA GCTGTGGCAA TTTCACCCTT ATTCCTTGGA GAGGCCAGCT GCCCTGCTGGA AGGAGTCAGA AGTCGGTGGA TGTCATTGAG GCCTTGGAGG CCCCAGTNTG GCGGGAGAGA AATCCACACC TGTGCCTGGA GTTCTCCTTC CCTGACCCTC TGAACCGGCG CTTAAAATGC TGTCCCGCCT GGAACAGGGA GGCCACATCC AGCAGTGCGT CCTCAATGTG CTGCCCCAGC CTGTGGGAAT CCGTTTTTGT GCTTGATTTT TTGCTGGAGA TGTGGAAGGT GATCATGCCA TCCCCCATGA AGATATAAGA AACANCATAA CCATGGTCAT CAGCAGG

SEO ID NO:1995: (Length of Sequence = 341 Nucleotides)

GEACCTATAT GEOCATECTO TEGOTOTACO CITGEGAACO CIGATOCCEG TETETCECCO ASCITETTAN ESPECTEGGA
TECTECATOT COAGGOAACT ATGOACTITO COGGGGAGA AACCAGTATG AGAAGTGGGG GCAGGGCACA CATTOATOTT
TETACOTGCO TOTTTGGTTT GGACCTGGCO AGTOGGGTCA CTGCCTCCAC GTCTGAGGCO COGCCAGCTG GCCGTCTGTC

CICGCCAGCC TCAGGCTGCT GCGCTCTCTC GGCTTTTACG GACCTCTGAG GCCGAAACCC CACCTCGAAG TTTCCCCGTG ACAGTGCGTC CGAGTCCACA T

SEO ID NO:1996: (Length of Sequence = 316 Nucleotides)

GCATATGGTT GGTGAACAGT TTTGCAGCCC TAGGCTCCTG TACTGTGCGT GCACCGCCGC CCGGGCAGCC GCTGGCTCCA GCTCACGAAA CAGCCCCGGG CGCCGCGCCG CTCTGAGTCC AGCCTCCTAC TGAGAACAGT CCCTCCCTTG TGCGGGTCGC ACGGCTAGCC GCAGGTTCGG CCACGTCAAA TCCATTTTNT AAAAAAGCAG GGAGCAGAGC TCTCTCTTCG CCGCCGACGC AGAAAGGAGC TNGGGAGGAA AAAGCTGCTG CCTTTTGCGC TGGAGATTCG TGGGCAAGGC TTCTCATTTT CCCAGG

SEO ID NO:1997: (Length of Sequence = 320 Nucleotides)

GCAGGITIAT GITTITATIT ATGIATITIA ACTGACTIAT TIGIGIATIC CACTAGAACA ATACATICAC AATATACITG CAGAACIGIG CCTGGNGCAT CATGGGAGCA GAGAACTIGI CCAGTGAATA GITGITGAAG AAAGGNGTAA AATCICCCCC AAACCCTAAA GGCATCCTIT TCGTAGTGIG TGTCCCATAG GTATGGCTGC TGAGCACCAG GGCTGCTCAC CATGCTCCCA AGAAGCAGAG TCAGGGAGCA AGACAGCAGG GTTTATTAAG GTGCACANCC ATGICTGAGC CCCAGCTCTC TCCGNCTTCT

SEO ID NO:1998: (Length of Sequence = 395 Nucleotides)

TTTGATGCTA TGGCGCTGGA CCCAGGGCCC TCCCAGGCCA TCTCTGTTCC TCTGGGGTGG TCCAGTTCTA GAGTGGGAGA
AAGGGAGTCA GGCGCATTGG GAATCGTGGT TCCAGTCTGG TTGCAGAATC TGCACATTTG CCAAGAAATT TTCCCTGTTT
GGAAAGTTTG CCCCAGCTTT CCCGGGCACA CCACCTTTTG TCCCAAGTGT CTGCCGGTCG ACCAATCTGC CTGCCACACA
TTGACCAAGC CAGACCCGGT TCACCCAGCT CGAGGATCCC AGGTTGAAGA GTGGCCCCTT GAGGCCCTGG AAAGACCAAT
CACTGGACTT CTTCCCTTGA GAGTCAGAGG TCANCCGTGA TTCTGCCTGC AACTTATCAT TGATCTGCAG TGATT

SEQ ID NO:1999: (Length of Sequence = 337 Nucleotides)

GAAAGTATIT GIGIGATIGA GICACACGCI GAATCAATCI TCATATAAIG CCATITITGC TIAAAAGAAI GCCAGACITG
GGCATTAGGC TGACATITIC TIGAAAACAG TGAGGCITIG CITTAGGGAA AATAGIGGIA GIATITATGG TCGATGATAA
AGITCCIAGA TITTAAGCAA AAATTITAGA AAGCITGIAT CAGCIGCIGI AAGIATATAA TGAAATCIGI CATTATITGA
TTATCIGCAT AACIGAGICA GIATITCCAA AIGATCAAIG CATAGIATTA TAAAAATCAT ACATGGGIAA GAAATCITTA
CAAAGIGICA GCTAGAC

SEQ ID NO:2000: (Length of Sequence = 329 Nucleotides)

ATGTAGCCCC CTGCTGCAAA GGTGCCATCT TTTTNCTGCT GCTCACACAG CAGCGTGCTC AGGGCCTGCC TGCATGGCAG
NNTCATCATG GGGAAGCCCA CAGCCACTGA CATCATGAAG CCCACACGGA GCATCTCCGT CACCAGGTTG GAGGGAAAGT
GCATGAGCAC GTTTGCCGGC CGTGGCCTCG GTGAAGCTGA CGTAGCCGAA AAACCCCACC ATGACGTAGG AAGGTGGTGA
CCACATTAAG GGAGGAAGCA AATATGGAGC TCATGGTTTT CACTTGACGG GCTCATCCAG GCTGTCGTAG GTGGGCAGCA
CCTGGGACT

SEO ID NO:2001: (Length of Sequence = 308 Nucleotides)

AAGTCTGGGG TTTGGTAGGC TCCCAGGATT TCCCTCAGCA GGCATTTGTG CTGCCGCAGG GCCGTCTGGG TGCCCCGCAG GTCNTCCTGG ATGCTCTGTA GCCTCCGGTX GAACGACTCC CTCACTGACT GTGTGGCAAA GCTGAGCTCT GCCCTGACCC ATGTGGCATT GGCCAGGATG GGGGCCANGC CCTGTGGGAT GCTTTGCTGC CCGTNTCCTG AGGCACCGAC TGCCTCTCCT CCCAGTGTCC CCAAGTGCTT CCTCAGAGAC TCAACCTGCN TCCAGAACTC ACCATCCACT AGGACCTT

429

PCT/US93/01294

SEO ID NO:2002: (Length of Sequence = 242 Nucleotides)

AGCCAGGCCC TGGGCCCAAG CCCCTTGTCC CTTCTCCACT GCCCCTCTTT CCAGACAGTA AAGGCCATGG TCAGTGTGTT
TTTCTCTTGT AAACAAACCC CAGCTTGTTT AACAGAAATG CTAATAACCT ACTGGGAAAG ATGGAGGTCT AAATTACCTC
CAGGGTTTTT CTGGGGGTTT ATCACCAGTG TGGGTCCCTT CTGATACCAC CAGGTTCACT CCAGGCAGAG TGGGGCGGAA
GG

SEO ID NO:2003: (Length of Sequence = 328 Nucleotides)

ATATTCTCAC TTATAAGTGG GAGCTAAATN ATGGGAACAC ATGGACGCAT AGAAGGGNAC ACTITTACAC TNCTGGTGGG
NGTGTAAACT AATACAACCA CTGTGGAAAA CAGTGTGGCG NTTCGTTAAA GAACTAAAAG TAGATCTCCC GNTTGATCCA
GCAATCCCAC TACTGGGTAT CTACCCNNAA GAAAATAAGT CATTATACAA AAAAGATACT TGCACACACG TTTATAGCAG
CACAATTTGC AATTGCAAAA AATATGGGGC CAACCCAAAT GCCCATCAAT CAATGAGTGG ATAAAGGAAA TGTGAGATAT
ATATATAT

SEO ID NO:2004: (Length of Sequence = 211 Nucleotides)

AGCCTTTTTA TTATTGTNIT TTTTTTTTT TAANCGAAGG TCCCTTACTG GTCCTGCTTC CATGAGTAGC CGTGACCAGG GGAAAAGGGA GAGGAACCAG CCGGCACAGG GAGGGGTCAT CTCCACAACA TTCCATTTAT ACACAGAACT AAACAGACAA GCACAGNGTC ACTATTGCGG TTAGAAGTTG GCAGCATGGG AAGGGGGAGG A

SEO ID NO:2005: (Length of Sequence = 241 Nucleotides)

SEQ ID NO:2006: (Length of Sequence = 266 Nucleotides)

TTCCCCCTAA CCTTGTGAGT GGGCCTTTTA AGTAGTAAGT AGTATACACC TAGATATGGA TAGATAGCTA GGTGACCAAA CCTAATGGAT TAAGGCCATC CTCGCCTAGG TCACTTACTA AAGATCAGGT CATATGTCAT ATCGTTCCTG TGCTTTTTAG AACGTATTTG GGAATGGGT CCAGATTTTT TTTAAACACA TATTAAAGAT TATTTATATT ATGCTTTGTT TCCGAAAGGT TTTAAGGTGG ATTAAAATAT AAGATT

SEO ID NO:2007: (Length of Sequence = 419 Nucleotides)

AGAAAGAGGC TITCTITCIGC GGAGGCAGGI GGAGCACAGG GAGGGCTCCT GGGAGGCACA GGAGTGGGGI GGGGCCCACGG
AAGGGGGAGG TGGACAGAGC GACTIGGATA AGGCTGGGCC GGGCCCACGC CCACCTCAAG AGGGGGGCCG CCTCCTCAGG
AGGNATCAAG GTGCAATCCA GTCTTCCTTT CTCTCCCTGA AGACCTGAGT TCCAGCCTTC ACAGAGCGTC ATGCGCATTC
TTCTTTCTGG ATGCTAACCC CAAATCCGAC ACTCAATGGI GCACCTCAGG TACCTGCCAA GGNTCINTGG GCCCACATGG
AAGGTGCAGG GTCTGGGTCC CTGGATGACG AGGTGAGGGG CAGATGGGTG ACCAGGGAAG GGCATGACCC AGAGCTNCCG
GGACTCATGG AGGATTNGG

SEO ID NO: 2008: (Length of Sequence = 360 Nucleotides)

CTTTTCTGGA GAAAATAATA CGCTCGTTCC TCTAATTAGC CCATCGGTTT CAGGITCATC ACTCTGCTAT CTTCTCCTGG
AGTTTACACA AGCCCTTCAG AGTGTAAACA CCGATGTGGA TTCAATCCCA CTCATTATTT TTTTCAATAA AAAGAGAACT
GTTTCAACAG ACAGGTGTG TTTCCGACAT CATCAGAGAG GAAGGTGGAT GGTTCTATAC GGTAAGCATT CTACCCTTCA

المهاج فوالوشاؤنان

GCTGCCAGGG ACAGATCCAT AAAANTCCAA AAAGGGAAGA GAGAAACAGC TTGAGTACAG CTGAATCATT CACAACAATA TTACAAGCAA TTACTTCAAT GGTAAAGTCT CCAGTCTAGA

SEQ ID NO:2009: (Length of Sequence = 411 Nucleotides)

ATTACGGGCA CCTGCCACCA CGCCTGGCTA GITTITGTAT TITTAGTAGA GACGATGITI CACCATGITG ACCAGGCTGG
TCTCGAACTC TIGACCTCAA GTGATCCACT CGCTTCGGCC TCCCAAAGTG CTGGGATTAT AGGCGTGAGC ACCTGTGCCC
AGCCTCACAG CTGCATCTTA ACCTTACCTT TGCCTCTGCC TCTCAAGCTG GTACCTCCTA ATTTACATCC TAAGAGTGGA
ACCATGTGAC AAGGACTGGA GTGCCATTGG CTGTGGACTG TTCAGGCAGG GAAGTACAAG ACCACTCTTG TATTCAGGGG
CAACCAAAGG AGAGAATTAC GTACTTGTTG AGTACAAACT GCACCAAGCC CTGGAGACCC ATTACCACCG TTAACCCTCA
ATTACAGCTCT G

SEO ID NO:2010: (Length of Sequence = 311 Nucleotides)

AAGAAAGATG CCAGCTCTTT ATTACCAGGG AAGCTGTGTG CACGCGCGTG GAGGGINCCT NTGGAGCTGA CCGGGCCCTT
ACCTTCTCCT GCTTGTCAGA GGTGAGTCCT GGTACCCAGC ACGGTGGCCT CCGGGAGGCT TTGATAGGTC AGCCTTTGCT
GCCTCCCAGC TCAGGGCTCC TCCAAGGAAC CTGCGGGGCC CCATGTGCCC ACAGCCCGAG GAGGGAAGCA CCGACCGCCC
TCCTCGTGGC CAGTTGACAC ATCATCCATT TATTATCCTT CAGAGTCTAA AACTTCCTCG TGATACAACG T

SEO ID NO:2011: (Length of Sequence = 192 Nucleotides)

TCAGGACATT TCAGTGAGGC CACCTACAAG CAGAAAGGAG GCCCAGGGCT AGGGACAGAN TGGCCCCAGA GCCAGTCAGC TGCAGCAATT CTTGTGAGAA AGGGAGGGCA AGCTGCCAGA GCANTGINGC CCAATATGAT GCCTACACGA GACAGATGTC CCCAGTAGAG TGTGTTCAGT GACCTTCTAA AC

SEO ID NO:2012: (Length of Sequence = 367 Nucleotides)

GGATGACCTT CGAGGACGTG TGCCGGTACT TCACGGACAT CATCAAGTGC CGCGTGATCA ACACATCCCA CCTGAGCATC CACAAGACGT GGGAGGAGGC CCGGCTGCAT GGCGCCTGGA CGCTGCATGA GGACCCGCGA CAGAACCGCG GTGGCGGCTG CATCAACCAC AAGGACACCT TCTTCCAGAA CCCACAGTAC ATCTTCGAAG TCAAGAAGCC AGAAGATGAA GTCCTGATCT GCATCCAGCA GCGCCAAAG CCGCCAAAG CCGCGAGGG CAAGGGTGAG AACCTGGACA TTGGCTTTGA CATCTACAAG GTGGAGGAGAA ACCGCCAGTA CCGCATGCAC AGCCTTCAGC ACAAGGC

SEO ID NO:2013: (Length of Sequence = 213 Nucleotides)

GATTITATES AAAAAAATTI CCATTITINIT TAAGAAATAA GGAGTITINIG TGTCGAGGGC ATGACTACGA GAGGCTGGAA
GCTTCCAACA GAGAATGCTG AACGANITCC CCCATGCCAT CGCCATGCAG CACGNCAACC AGCCCGATGA GACCATCTTC
CAGGCAGAAG CTCAGTATTT GCAGATATAT GCTGTGACTC CCATTCCAGA GAG

SEO ID NO:2014: (Length of Sequence = 333 Nucleotides)

GITAAATAAA ACAGCAAATT CTTAAATACA TTATGAGTAA AGAAAGATTA AAATAAGGNA ACAGTACTTA CTGTGCAACT
TTAAATTATA CCAAGTAAAG TACACCACCT ATTCACTGAT AACATTTTCC CTACGTTGAA AACACAAAAC CTACTTATCG
ATATTTTTGA TATTAAAAAA AAGGACATTC ACTATTGTAG CCCTGACAAC TCTTCCAGTA TTTTTAACCA TTCAGATGTA
TTATGTGGGN ATATTTATTA ACATAATTIN GITTAACACA TTTCTTTCTA CACAAACTGA AT.TTAAAAG TGTCTATAAC
ATTTTCAATT ACA

SEO ID NO:2015: (Length of Sequence = 179 Nucleotides)

NCACCACTTA TIGICITCAA ACATTATIGC ACITTAACIT TCTTAATTIG ACAAAGCATT CAAGAAACAT CIGCAGACTA GTTTTAACAG ACAAATAACA CCTGTAAGCA GACATGACIG TCCTAAATIG TITATTAAGA AAGITAAAGN GCAATAATGT TIGAAGACAA TAAGIGGIG

SEO ID NO:2016: (Length of Sequence = 293 Nucleotides)

TTTTCCCTCC CCAGAGATGC TTTATTACAT GGTTTCATCA GTCATCAATG ATGGGTCCCT ATGCCCATGC GAGGAGACAG
GAACATCTGT GTGGGTACATG GCACTGTTCC CCTCTCAGCT ACGCAGTCAG ATGGGGGCAG GGGGATGAAT GGGTGCTTGG
CTTCCCTGCT GTTGGGCAGG CTCTGAGATC TCAGCAGACA GAAATGAAAG CCTGGCAAAT AGGGAGGCAG GAATGTTCAA
GCATCGGTGA CCTCCATGTT CTGCAGCCTG TTTTCTAGGG TGACGTCTCT TTG

SEO ID NO:2017: (Length of Sequence = 504 Nucleotides)

CECETECTES COECECTETE GEOCOCCICE TIMECONCCC CAGNOTOCTIC GIOGOCCICES ATATOTETIC CAAAAACCCC
TECCACAACG GIGGITTATE CEAGGAGAIT TOCCAAGAAG TECGAGGAGA TETCTTCCCC TOGTACACCT GCACGTGCCT
TAAGGGCTAC GOGGGCAACC ACTGTGAGAC GAAATGTGTC GAGCCACTGG GCATGGAGAA TEGGAACATT GCCAACTCAC
AGATCGCCGC CTCATCTGTG CGTGTGACCT TCTTNGGNTT GCAGCATTGG GTCCCGGAGC TEGCCCGCCT GAACCGCGCA
GGCATGGTCA ATGCTGGACA ACCAGCATCA ATGACGATAA CCCCTGGTTC CAGGTGAAAT TNCTNCGGAG GGATNTGGGT
AACANNINIT GTTACGAAGG GTGCCANCCG TTTGGCCCAGT ATTGGTACCT AAAGGCTTTA AAGGTGGCCT ANAGCTTAAT
TGGNAGGATN CGNTTTNTCC ATGT

SEO ID NO:2018: (Length of Sequence = 354 Nucleotides)

AGANCAGACC CACAGGCATG CAGAAAGGTA GGGCAGTATG TITAANICCA GACTIGGCAC ATGGCTAGGG ATACTGCTCA
CTAGCTGTGG AGGTCCTCAG GAGTGGAGAG AATGAGTAGG AGGGCAGAAG CTTCCATTIT TITCCTTCCT AAGACCCTGT
TATTTGTNIT ATTTCCTGCC TTTCCGAGTC CTGCAGTGGG CTGCCCTGTA CCCTGAACCT CATGAGCCTC TAAGGGAAAG
GAGGAACAAT TAGGACGTGG CAATGAGACC TGGCAGGGCA GAGTACAAGC CCAGCACCCA GTGTCCCAGN CTTACTGGGT
CCTTANCCTG GGCCAAACAG GGAGGGCTGA TACC

SEO ID NO: 2019: (Length of Sequence = 295 Nucleotides)

GACACAACCT TITGAACTAT TECTECTETT TICATITIAA AAAGGAACTT TITATACTAA AATTATAGGA AGAACATAAT
ATCIGACGIC ACGTAAATTC AGATTTGAAG GAAATTTACT TITTINCCTT ATTIGINCIT ATTITITCCIC ATTITGITAA
GAACCAGCGA ACACTITGAA GAAAGCCAAA AGITTACATC TGGAGCTGGA GGGTTCTGTG ACTGCACACC AGGCACTCTG
CCAGCCCTAC TICTGCCTGT AGTCCTGCAG GTCACTTGCC AGAGGTGGTA CTTTC

SEO ID NO:2020: (Length of Sequence = 217 Nucleotides)

ATTGGAACIT AAGTITCACA AGGAAAGIGG TCACTITAGT TCACCACTIT CCTTGTGAAA CTTAAGTTCC AATGGGAGAA TGACAGTAAA CAGACAACTA TTATAATANG TCCATGGAAG ATTITGGTGT ATGINAGAIT TNCAAATCTG TAGAGAAACN TNGGCTCATT CAATAAAAAT TTTGAAACCA TTGATTAATG TCTTAATAAC TATATGT

SEO ID NO:2021: (Length of Sequence = 380 Nucleotides)

TITITICITÀ AAACAACAGC AACGIGATCI TGGCIGICIG ICAIGIGITE AAGICCAIGG TIGGGICITG TGAAGICIGA GGITIAACAG TITGITGICC IGGNGGGATI TICITACAGC GAACACTIGA GITCCICCAA GICCCAGAACACTIGA GGCAAGAAGG ATCAGGICAG CCACTCCCIG GAGACACAGC CTICIGGCIG GGGACIGACI TGGCCATGIT CTCAGCIGAG CCACGCGGCT NGTAGTGCAG CCTTCTGTGA CCCCGCTNTG GTAAGTCCAG CCTTTCCAGG GCTGCTGTGTGAGAGTGCAG TCTTATCGAG ACCCAACGGC TCAATCTGCT CATCCNTAAA GTGGGGGATA

SEQ ID NO: 2022: (Length of Sequence = 223 Nucleotides)

GGTCACACAG CTAGTTGGTA GAGGAGCTCT TCCATAAGAT AGCAAGGCCA CATCACCTGC AGGGCAGTGC CTGCNCTGGG AGGTGGCACA ATGTGCCAAG TGATGACGAT GACAATAACT ATGAAAGGAT TTTATATTTG CACAGCATTT GGTTGCCTGA TCTTCGATGA GGAAGAGCTC CTGCCGATGT CTGCTGAATT GTGCAGTAAA ATATTCAGGA TGG

SEQ ID NO:2023: (Length of Sequence = 294 Nucleotides)

TATTCTTAAG TITGCACTIT ACAAAACCAC AAGGGAGAAG TCCTTGAAGG GGAGACAGGG GTAGGGGATT AGGGAGTGGG
GGATGGTAAA GAGGGGAAGA GGAAGACCCA GAAACGAAGT CCCCTCCAAC CCCATCTCGG GGACCAAGCA GAGACTAGGC
CTCAGGCTAG CCCAGCAGGG TTCCTGTGTC CTGGTTGTAC AGAGCTAGGC CAAAAGACCT CAGGGGAAGG GCCATGGCCC
TCTAGAGACT GCCGCCATTT GAGGGACAGC CACAGGCCAA TGTTTCCTGT GCCC

SEQ ID NO: 2024: (Length of Sequence = 234 Nucleotides)

ATTITICICE GEITGCAAAC GICTICCTGC CITGAGCTGG GAGCTTCACC AGGCTTCGGT GTAGCGGACG TCCACITCCT TCAAATTGGG AAGCTTGGCC TTCAGATCTT CGTAGGTGTC AGCTGAGAGC TTNGTGCTGT TCATGTTTAA ACTGCAGAGA CTCTTCATGG AGCTCAGGGC CAGCAGGCCA GCGTCTGTAA CCGGGGTCTC GCACAGGTTC AGCACCTGGA GCAT

SEQ ID NO: 2025: (Length of Sequence = 327 Nucleotides)

AGGAACAAAT GTTAAAGGGT AAGATAATTT CCCTGCAAAA GGACACAGAA GGCAGTCTTA AGAAGATGAA TGGATGAGAG AAGGGAGAGA ATAAAATGCA ATAACGAGCC AGCATTTACT ATGTATTTNN TCCTCACCTG TCTCTCCATA TTTAGGTCAC TTACCAGTTT CTGTGCCCTT TTGGAGCTTT TNTTGAGGGC TTCATTCTCA CCCTGTATTT CTTTAGCCCT AAATTGACAC TCTCTCCAAA AATCCATTCC ATTGTCTGTG GACCNAGATG TTCTATGTAA TTCAGAAGCA GAACTCTTGG CTAAAGGGCT AGTGTGG

SEQ ID NO:2026: (Length of Sequence = 328 Nucleotides)

TCAGTATAAA TITAAAAGAA ACAGCITAAT GAAATACAAG TCAGTITATT TGATATTCAG CCTACAGCIT TCCAAAGCAG CAGTIGAACA TGITGITGAG TITATACCAT TCATTCATTC ATTTATTTTT NCTITCTITC TTTCAGAAAA TACTGGGTGT TTGATATTTG TTTCACTGTG CTAGTTTCTG GGAATGTGTA AGGAAGAGGC TGGCTGTGTG GATGAGAGCA ACTTGCTTTT TACAATAATT ATTTGTTATT GTAAATTAAC AATTTGCTCT TCTGGTATTA TATGGAAGTA TTTGATCCNG TTGATGGCAC TGCCTTTG

SEQ ID NO:2027: (Length of Sequence = 307 Nucleotides)

AAGAAAGATG CCAGCTCTTT ATTACCAGGG AAGCTGTGTG CACGCGCGTG GAGGGTNCCN TTGGAGCTGA CCGGGCCCTT
ACCTTCTCCT GCTTGTCAGA GGTGAGTCCT GGTACCCAGC ACGGTGGCCT CCGGGAGGCT TTGATAGGTC AGCCTTTGCT
GCCTCCCAGC TCAGGGCTCC TCCAAGGAAC CTGCGGGGCC CCATGTGCCC ACAGCCCGAG GAGGGAAGCA CCGACCGNCC
TCCTCGTGGC CAGTTGACAC ATCATCCATT TATTATCCTT CAGAGTCTAA AACTTTCCTC GTGATAC

SEQ ID NO:2028: (Length of Sequence = 272 Nucleotides)

ATCCATTACT GCATTAACCT AGAGTTAAAA AGGAATATTG TTTATTGTTT GGCTCTCCCC ACTAGAAGTT TCACAGGNGC ACAGATCATA TCTACCATTT GAACAGCTCT CTGCCTGATG GCTAATACAT TINITGGCAT ATAGTAGGTA GGTGCTCAAT

AAATTINITA CAGGAATAAA TGAGATAGGA TITICAAGGG TATTINCTAT TAGGATTTAA TAAAACAAAG TGATCITIAG AGAAACAAAT CICCCCATCA ACATGCIATA CT

SEO ID NO:2029: (Length of Sequence = 261 Nucleotides)

SEO ID NO:2030: (Length of Sequence = 384 Nucleotides)

NNCCNNGGAC CAACAGCAGC CAGAGCAGTT AGCCAGTTAG TCCCCAGGCC TGTGGCACAG GCGTTTCTGA CCTGCTGGGC CGAGAATGGG TAAGTTGTCT GGAGTCAGGT GGGCCCACGT AGGACAGGGT CACAAAGCCT GGGTTTGTTT CTGGGTACTT TGCGCCCTCTG GGGTGCTAGA GGTGGGGGCAT GGTGGCTGGA AGTAAAACTG CCAACTCTGG CCCTCAGAAC TCTCAGGTAT AGAAGCCCAA GATGTCTAAT ACCCTNTCCC AGTGCCCGAG AGCTGCCTGG TGTCAGGTAG AGAGGACACT GTACCTGGGT GAATGATCAG ACCCTGGTAG ACCTGGTAG ACCTGGTAG ACCCTGGTAG ACCCTGGTAG ACCCTGGTAG ACCTGGTAG ACCCTGGTAG ACCTGGTAG ACCTGGTAG ACCCTGGTAG ACCCTGGTAG ACCCTGGTAG ACCTGGTAG ACCTGGTAG ACCCT

SEO ID NO:2031: (Length of Sequence = 261 Nucleotides)

ATCACAGAGG AGAAGCCACT GITGCCAGGA CAGACGCCTG AGGCGGCCAA GGAGGCTGAG TTAGCTGCCC GANTCCTCCT
GGACCAGGGA CAGACTCACT CTGTGGAGAC ACCATACGGC TCINICACTT TCACTGTCTA TGGCACCCCC AAACCCAAAC
GCCCAGCGAT CCTTACCTAC CACGATGTGG GACTCAACTA TAAATCTTGC TTCCAGCCAC TGTTTCAGTT CGAGGACATG
CAGGAAATCA TTCAGAACTT T

SEQ ID NO:2032: (Length of Sequence = 344 Nucleotides)

CCCCTGCACG GCGTCTGGTT CTTCGGGGAA AACGCTCACC CACCCCTGGT AAAGGGCCTG CAGATCGAGC ATCCCGGGCC CACCCTCGAC CACGCAGCAC CACAAGGCCAG GTCACCCCAG CAGAGGAAAA GGATGGACAC AGCCCCATGT CCAAAGGCCT AGTCAATGGA CTCAAGGCCAG GACCAATGGC CTTGAGTTCC AAGGGCAGCT CTGGTGCCCC TGTATATGTG GNTCTCGCCT ACATCCCGAA TCATTGCAGT GGCAAGACTG CTGACCTTGA CTTCTTCCGT CGAGTGCGTG CATCCTACTA TGTGGTCAGT GGGAATGACC CTGCCAATGG CGAG

SEO ID NO:2033: (Length of Sequence = 373 Nucleotides)

GGAAGAAGA AAGAAAGAAA GAAAGAAAGA AAGAAAATGG CCCCATAGTG CTTAAGTCCT CAGACATGTG TCCTGGTGCT
GGGGACAGGG CTTCTGACAT TCTCTCAGGT CAGTATTTGC AGGTCATCCA CCTTCGACTT CAACACATGT GACCAGAAAC
CTTCCCAAGG CAGCCATCCA CTTTGCTGTC CCTCCGACGG CCATGGCTGA CCACTGCTGC TGCTGTGTAT CCTCGGTGAC
ATCTGGCCTT GGCAGCCTAT GGATTINTGC CATTCTCCTG GCATGAAATC ACTCCTTCTT GTTGTTTTAA TTTGCATTTC
TTCAGTTACC AGGGCAGTTG AGCATCTTTT CATACACTTA CTGACCATTT CTA

SEO ID NO:2034: (Length of Sequence = 289 Mucleotides)

CCACCAAAGA ACATCACGCT GICTTATGTC AAATGCTCGA CAATACCTCT CAGTAGGACG TIGTTGCAAG GCTAGCTAAT
TITAAATCIG GIATGAGTAA TACAGTCAAA CCTAGTTAGT ATGCGAGAAA GTCGTTGCTA ACGCATGGIG AGAGGATGTG
ACGTCACAGC ATGAGCAGTC CCTGGTTGTC CCATTGTCAG ATAAACGTAG TANAGGTAGH CCAAGGTTCT ATGCCAGGTC
TCTGAACCCC AAAGCCAGGC CTTTCACTTT TGCTGGGTGG CCTGGAAGC

SEO ID NO: 2035: (Length of Sequence = 290 Nucleotides)

CTITICCTIC ATCTGAACAC AGAAGGAGCC ACGITCTGGA AAGINTGCCT GTCCTTCCCG GGAGTGGGGA GGCCGGTGTG
AGTITTGATC TTCCAGCTCA GGCAGACACC TTACACAGTG CAAACAAGAG CCGTGTCAAG ATGAGAGGGA AGCGTAGACC
GCAGACCCGT GCAGCTAGGC GGCTGGCTGC TCAGGAGTCC AGCGAGGCTG AGGACATGAG CGTCCCCAGA GGACCCATTT
GCACANTGGG CTGATGGCGC CATTTCCCCA AATNGCCATC GGCACCAGGT

SEO ID NO:2036: (Length of Sequence = 241 Nucleotides)

TTATTTTATA TAAAAAGIGT TTCTGTGATT CTCCAGAGCC CAGGAGTCAG TNCTGGTGGT TGGAGGGACC TGCCCCCACT
GGTTCATTTA ACCCTCTGTC TCGGTGCCCT NAGAACCTCA GCCAGAAAGG CAAGGAGGAA ATCAGAGCAN GAGCCTCATA
CTCTTGGTGA TCTATTCATT CTNTGACCTC AGGGGTCACA TATAAGGTCA GTGTTTCTCG TCCCCGNCGG ATCTGCACTG
C

SEO ID NO:2037: (Length of Sequence = 270 Nucleotides)

CTATTATTT GCATTITTEG TAGAAGGGGT GGTCTCACCA TGTCGCCCAG GCCGGTCTCG AACTCCTGAG CTCAAGCGGT
CCACCTGCCT CAGCCTCCCA AAGTGCTGCG ATTACAGGCT TGAGCCACTG CACCCTGCCC AACCTTGACT ACTTCTAATA
GGGATGAGTC GAGTAGCAGT TNGGGGCGTC CTGTGCGGCT GGGTCTGCCT GAGGCTCCCC TCGGCCCCGT CCATGGCTTG
TTGTGCATCT GGCCCTGAGT GCCTTGGCCC

SEQ ID NO:2038: (Length of Sequence = 151 Nucleotides)

SEO ID NO:2039: (Length of Sequence = 166 Nucleotides)

TITIGICIETT ACAACCICCE TATEACECCA CECCACCCEC TETTCACETC CCGTCGECCI CCTGCCACAGN CCACACECTG CCCCCGGAAG GCCCCTCCTG TEGAGAAGCC GGACCCATCC CCGAGGTCCC CAGCGAGGAC ACANACTCCA CGAGAGCAGC CCCTCC

SEO ID NO: 2040: (Length of Sequence = 362 Nucleotides)

GAAGTACGGT TAAAATTAGA TITGACCATA TGGAAGATCT TITTACCAGIT GGTCTCCAAG AATGTCTTCC TTATTATGTT
ATTGGTCATT TITGAGCGTG TGTGTTGGTG GGGTGGTTTC TGCCTTATAT TCCTTAACTA CATTGTATAT TTTTGTAAGG
AATTGGGAAT TCATTTTAAT GCTTTTTAAC ATCTTCACTG GGAACTGGAA TAAAGTTATT CTTGACTCTG TACCTTGAGC
CATTGTCAAA GTCAGGGGTT ACATTTTAGG TATCTAAAAA TTACTCTTTA ACTTTCACAT TCCCTGGGTT AGGAAGCTGC
TGTTCAGGAG AAATTTTCCN GGTTCTTCTG GCAATTGGCT TA

SEO ID NO: 2041: (Length of Sequence = 360 Nucleotides)

SEO ID NO: 2042: (Length of Sequence = 403 Nucleotides)

ومضورة كرواء

GTTATTGTTG TTTGAGATGG AGTTTCACTT TTNTTGCCCA GGCTGGAGTG CAGTAGCATG ATCTCAGCTC ACTGCAACCT CTGCCTCCCG GGCCCAAGAG ATTCTCCTCC CTCAGCCTCC TGAATAGCTG GGACTACAGG TGCCCAACAG CACACCCGGC CAATTGTTGT ATTTCTAGTA GAGATGGGGC TTCTTCACGT TGGCCAGGCT GGTCTCGAAC TCCTGACCCC AGGCGATTCC CCCACCTCAG CCTCCAAAAG CGCTAGGACC ACAGGCGTGA ACCACTGCGC CCAGTCGGAA GTAATAGTTA TTAACCAATG TGATGGCCGG GTGTAGGGAC CCTCGCCTGT AATCCCAGCA CTTTGGGAGG CCAAGGAGGG AGGACCGCCC GNGACCAAGA GTT

SEO ID NO:2043: (Length of Sequence = 331 Nucleotides)

CCCCGTACGE TGTGGCTCTC AGCAGCCTCA CCACAGGCAC CGCAGCTTTC CCGCTGTGCA CCCAGCTGGG TGTGTGAATC CCCCTGGACT GCGCCCAGGC CACCTTCATC TCCCATGACA AGATGGTCAT CTCCCTCAAG GGCAGTCAGA TCTACATGCT GACCCTCATC ACCGATGGCA TGCGTAGGTT CCGAGTGTTC CACTTTTGAC AAGGCGGCCA CCAGCGTCCT CACCACCAGC ATGGTCACCA TGGAGCCTGG GTACCTGTTC CTGAGTTCTT GCCTGGGCAA NTCTCTCCTC CTCAAGTACA CCGAGAAGCT TCAGGAGCCC C

SEO ID NO:2044: (Length of Sequence = 244 Nucleotides)

ATGGAAGATA CTAAGAGCCT CAGTCTGGAA GCATTTACCT AGGAAGCGCA TATAGACAGA GAAGATCAAG GACTGAGGCC TGAGACAGTC AGCACTTAAA GGGTGAGGGG AGAAGTGCCA AGGAGACAAG GTGAGAACAG CAGAAGAGTA GCCAAGGCCC AGGATGTTGC CACAGAAGCC AGGAGAGGTG AGCATGAAAA CAGAGGAGGA CCAGCTGCTG GGACAGAAGA GCCATATGGA AGAG

<u>SEO ID NO:2045:</u> (Length of Sequence = 333 Nucleotides)

GTCAGGGATT TGTCCATTCT GCTCTTGGCC TCTCCTGAGG CCTCATAATG GGAGACCAAA TCAAAAATGT CCCATGTCAC
TTGAGTGGGT ACACTGCCTA CAGAACCTTG AGGTTGACTC CTGCTTCAGT TCTCAGCTGT TTACCACAGC CCTCCAGGGT
CCAAAGATTG AGGAGCTTTC TCTTTCCTGG GAGGAACTGT CTCANATTTA GCTTGTGTGT GTTTTGGACA GAGGCTCCAC
AGCGGTGGCT CTTGAGGAAT CCTCACCAGT TTGTNCTCTT CCCTCTGACA AGCAGCACCT GAGCAGATGC TGAGGCAGTT
CATTAAACCA GGG

SEQ ID NO:2046: (Length of Sequence = 274 Nucleotides)

GCAGGITTAT GITITIATIT AIGIATINA ACIGACITAT TIGIGIATCC CACIAGAACA ATACATICAC AATATACITG
CAGAACTGIG CCIGGNGCAT CATGGGAGCA GAGAACTIGI CCAGIGAATA GITGITGAAG AAAGGAGTAA AATCICCCCC
AAACCCTAAA GGCATCCTIT TCGIAGIGIG TGICCCATAG GIATGGCIGC TGAGCACCAG GGCIGCTCAC CATGCICCCA
AGAAGCAGAG TCAGGGAGGC AGACAGCAGG GITT

SEQ ID NO:2047: (Length of Sequence = 327 Mucleotides)

GECCECCECA ACCIGAAATC AGAGCAGGCG TCCGTGGCGC TCAGGAACCT TGCTGAGCTT CGCCGATCTT TCATTGTTGC .

TCCTCTTT

TCCTCTTT

TTCTTCTGGA TCCCTCTTGC GCCTCGGCA GCTGGTGCCA TAAAGGCAGA CCCGCGGGCG CGCGCCGCCA ACCTGAAATC AGAGCAGGCG TCCGTGGCGC TCAGGAACCT TGCTGAGCTT CGCCGATCTT TCATTGTTGC .

TTCATTT

SEO ID NO:2048: (Length of Sequence = 241 Nucleotides)

ACTITISTICI TOTGATTITA GGACTOTIGO TIGGOCATGIG CITANGGITG COTOTOCTIC ATTINICACT GGATTINICAC TIGGATCAT GGAGATACAA AGOGAGCAGI TOTTGGITCAG AACCOTOCTIC TIGGITTACAT TIGGITTACAT TIGGITTACAT GGAGACAGIA GCAAGGCCAA GCIGATGGCT GCTTGTTTAG GAGGCCATCA GTTCCTTCCT GIGGAGAAGG G

SEQ ID NO:2049: (Length of Sequence = 269 Nucleotides)

ATTITIAGIA GAGACAGGI TICACCATGI TGGCCAGGI GGICTCAAAC TCCTGGCCTC AAGTGAGCCA CCTGCTITGG CCTCCCAAAG TGITGAGATT ACAGGIGAGA TATTCTATAT TCATGGATIG AAAGACTCAA TATTGITAAG ATGTCAGINC TTTCTAAAGN GATTITITAG ATGCAACACA ATTCCAATCA AAATCCCAGG NITTITTIGI AGCTATCAAT TGATAGATAT CAACAGCCAG CTGATTCTCA AATTTACGI

SEO ID NO:2050: (Length of Sequence = 170 Nucleotides)

TTTTGAAGAG AACGTCAGTT TAATAAAGCT AAATGGGGAG AATTGAAGTT TGCATTTGAC ATGGTATTAA ACAAAACCAA AGGGCTGAAA CTCATGTTTA GACAACACAG GTCACTAGTC ACTAGGCAAA GAAAACAGTC CACAGCAGGT GGCACAAATA ATTCCTATAC

SEO ID NO: 2051: (Length of Sequence = 262 Nucleotides)

CAGGGCACAC GCAGGACCAC TGTGGATTAG AAACCCACAC GTGTCACTCG CAACATTCCT CCCACATCCA CATCCACGAC GGAGCCAAAT CTCATTTGTC ACCCTCAGTC ACCACCCCAC AAGATGGAGC CGCTGGTTAC GACATGGATG ACAGGTGTCA TGCACAGGAA GAGAATTTNT CCCCGGATAC CCCTGAGGAC CAAGGACCAC CCCCAGGCTA GGGTGGGAGG ATTGAGAGCA GTGCAAGAAA CCAAGGAGGA TN

SEO ID NO: 2052: (Length of Sequence = 325 Nucleotides)

GAAAAAAGAT TGTTTGTTA GAAAAAGCAA AAACAAAAAA GCATTAGAAA GTGGGAGCCA CTGCACAGCA GTAGCCTAGA
GACTGGCTGC GATATGGTAG CTCTGCCTTG ATATCATCTT CGTGTCTTCA GGCATAGAGA AATGGCAGAG GAGCAGTAAG
ACCCCCACAGG AGATGGCCAG AGGNTCCACC ATCAGCCTTC TGGGGACTGA GGAGGTGATC TTAGTGGAAT TATTTTATAC
TCACCTCCCC CGGGGTTTAG TCCTTCCTCC AAACACTTAG TTCCAGGGCG CAGGAGACCT GTTACTAGCA CTGTATGTTT
CTTTG

SEQ ID NO: 2053: (Length of Sequence = 222 Nucleotides)

TITCAAAATT AGTCTTAAGA GIATAAGCTG TITTINAGGG CTGTAGCCAG ACTACATAAT GAGCGGTGAA AGCGGCTGCC TTCCCCTCTC CTGACACCAG CAAGGGGGAG GCACCATCAC CGGCCCTGCC CCATCATGCA TCCAATGATT ACTAGCACTA GANGCCAACG GCAAAGGNCC CCGCGCGCTT GCTCGTGTTT AATCCAGGTT AAGCTATACA CG

SEO ID NO: 2054: (Length of Sequence = 341 Nucleotides)

GIAAATTAAG AATATGGCCC CAGAGTTTG TITATCTGGG GTCTGAGCAT AGATTTTATA TICTCTGTTG CGTTTTTTAA
ATCTAACTTT CTGTCTCCAA TGGAGAGAG ACAGGGAGGA TACAGAAGTA TTGCAGCCCA GATCCCCTAT CAGGGGGACA
GCTGGTGGGC AAAGCAGCCA CCCCACAGCC TTGTGGCTAG AGTACAGTGG GGTGGACCCT CCAGCCCCAA TAGCCCTAGT
ACCCAGCTGG CAGGGTTGCC CACCCCTGCT GTCCACCTGC TCCATCCTCT AGGGGTTCCA CAGGCCCCTG ACCGCACAGG
GAGGCTGGGG CCAGCCTGGT C

SEO ID NO:2055: (Length of Sequence = 258 Nucleotides)

CIGCCICAGC CICCCAAGIA GCIGGCATIA CAGGCGCCCA CCACCACACC TGGCIAATIT TIGIATTITT AGIAGAGACG AGGITICACI AIGITGGCCA GGCIGGCICI GAATITCIGA CCITGIGATC CGCCIGCCIC GGCCICCCAA AGIGITGGGG ATIACAGGCG TGAGCACCAC GCCCGGCCAA CITGCITTIC TCIAATGGCI GGCGATGITA ATITITICAC TGGCTIATIT ACCGICICCI TCIGIGGA

SEQ ID NO: 2056: (Length of Sequence = 292 Nucleotides)

CTCTTGACTC CGAAGGCTGG TGACAGACAC ATAAGGCAGC TCAAACTCTT GCAACTTCCG TACAAAAGAA AAGGCTCCAT
CCTCTTTTTC TCGAACTAAG AATAGACTAA AGTATCCAAT CAAGTCATCT GGAAGATCCA GCTTTGCAGC TACAGCCTCC
AGGACATCCT CAGTCTGATC TGAAGTTAGC ACGTTGACCA GAACTTTCTG CCCGTTGCTG AGCAGCACTT CCAAGGACAC
TTCCTCTGTG GGGACCTGCT GTGTCTCCTG TTGTGCCCGA CGCAGGAAAC TG

SEO ID NO:2057: (Length of Sequence = 293 Nucleotides)

CCAAAAAACT TGGGIGCCIG AAGGIGGGGI TITGATCATG GCCAGGCITC AAATTTAGGI CAGGCICIGG TGGIACATCC
TTATATGCIT GGIGCTCAGC ACAGGICAGA ACACACAATA GACCCICAAT AAATATTTGC TGAATTTGAA CAATTCCIGT
AAAAATCICA TTAAGAGACA TCAGCITGGG ACACAGITCC TCICTTACIG TTCCTTCTCC CAGAAGCTCC TGGAATGAGC
AGGICIGGGG GCAGGGGGCA CACAGGGCIG CIGCTCAAAT CGGAGAATGG CAC

SEO ID NO:2058: (Length of Sequence = 172 Nucleotides)

CTTCTACAGT CAAGGAGCTC AAGCTCGCCG GGCGACCCTG CTCCTGCCTC CCACATTAAT GGCGGCATCC TCGGAGGATG ATATAGACCG GCGCCCCATC CGGAGAGTGC GCTCCAAGAG CGACANGCCG TACCTCGCAG AGGCCAGGTT CTCCTTTAAC CTGGGGGCAG CT

SEO ID NO:2059: (Length of Sequence = 245 Nucleotides)

SEQ ID NO: 2060: (Length of Sequence = 318 Nucleotides)

ATGCCCTGTT AAGGAGCTTG GGCTTGATCC TCTAGGCAGG GAGCCGTTGG AGGATTTAAG CCAGGGAGTG CTGCGGTTGG
TCACACTOGC CATTTATGTA GATCGTTTTG GCAGCCAGGG GAAGGATGGA TTTNAGGGGG ATGAGATTAG AAAGCTGGGA
TATGAGTTAG GAGGCTGAAA GATGGTTGAT AAAAATNATC GTTGGGCAGC CGAGATAACT GACTTCAAGG ACATATACTG
GACTTATAGC AGAGCCTGTT GAGTCTTGCT TTTGCACACA GTTCAAATAA TCACTTAGTC ATGTGGTTTA TCTTGCCA

SEO ID NO: 2061: (Length of Sequence = 331 Nucleotides)

AAAAATAAAA ATCTATAAAC TACGGATCAT AAGCAACTCC TGTTTCTGTG GGTTTCACCA CATTCTCCAG AAACTGAACT
TTTGCTCATA AAAATTACAT AGAATGTAAA CTAATTCATT TTTTAAAGTA AATGCAAAAC TAAGGGTTAC ACAAGCACTG
AGCATCAACA CTGACAGAAT ATTAATTCTG AAGCCCATTA ACTTTGACAA ACGTTTATTC ATCTTTGCCT TCTTGAAGCG
TGTGACTATC CCAGTTTTAC AGGAAAAGCT TAAACAGAAA AAGTTAAATA ATAATCTCAA GGTTAGNAAA CTAAGACATA
ATTTCTAGCT C

SEO ID NO: 2062: (Length of Sequence = 316 Nucleotides)

CTAAAATCAA CCACATAATT GGACATAAAA GAATCITCAG CAAATACAAA AGAACCAAAA TCATAACAAA CACACTCTAG GGCCACTGCA CAATAAAAAT ACAAGTCAAG ACTAAGAAGA TCACTCAAAA CAATGCAATT ACATGGAAAT TAAGCAACAT ACTCTCAAAA TGACTTTTGG GTAAATAATA AAATTAAGGC AGAAATAAAG AAGCTCTTTG AAACTAATGA GAAGAAAGAT ACAACGTATC AGAAACTCTG GGGTACAGCT AAGGCAGTGA TAAGAGGAAA ATTCTTAGCA CTAAATCCTC ACATTG

SEQ ID NO: 2063: (Length of Sequence = 312 Nucleotides)

ATCCATEGCT TTAGCAAGAT CCCAGTGTCG GAACTCTCCT AGCAACTTGT NTTCATCCAG TGATACTGGT TCTNTGGGGG
GCACTTACAG GCAGAAGTCC ATGCCCGAAG TGTTGGAGTG AGCCGTAGAT CCCCAGCCTC CACTGACAGG CAGAACACCC
AGTCAGATAT TGGTGGCAGC GGAAAATCCA CGCCTAGCTG GCAAAGAAGT GAGGATAGCA TTGCTGACCA GATGGCTTAC
AGTTATAGAG GACCTCAGGA TTTCAATTCT TTTGTCCTCG AGCAGCATGA ATATACAGAG CCAACATGCC AT

SEO ID NO: 2064: (Length of Sequence = 294 Nucleotides)

TACCTAAAGA ATCCTCAGAT GGGAGACCCA GCCAGTITIGG NICACAAATT AGCAGAAGTC AGCCAAAATA TAGAGAAACT
GCGAGTAGAG ACCCAGAAAT TTGAGGCCTG GCTGGCTGAG GTTGAAGGCC GGCTCCCAGC ACGCAGCGAG CAGGCGCCC
GGCAGAGCGG ACTGTACGAC AGCCAGAACC CACCCACAGT CAACAACTNC GNCCAGGACC GTGAGAGCCC AGATGGCAGT
TACACAGAGG AGCAGAGTCA GGAGAGTNAG ATGAAGGTGC TGGCCACGGA TTTT

SEQ ID NO:2065: (Length of Sequence = 331 Nucleotides)

GAGCIGAGIT TCACCGIGIT GCCCAGGCIG GICTCGAACT CCCGGICICA AGIGATCCIC CIACCICAGC CICCCAAAGC ACTGGGATTA CAGGIGIAAA TCACIGIGCC CAACCIGCIC AAACTCITGG AGAGAAGCAA GICTICIAGC TGAACGIGAT AATGGCCTCA AAAGCAGIGI TGACAGCAGA TAATCITCAC ACAGACAAAT GICTACAGIT TCTAAATAAG CCAACTGIGC ATATGGCCTA CAGGICCTIC AGCATAACCT ACCCAAAGCT CAGGITCCCT GAAGGCCAGG ACAGIACCTC GGGCCITCAA GCAGCATTIG G

SEO ID NO: 2066: (Length of Sequence = 321 Nucleotides)

GICITGANCT CCTGACCTCA GGIGATCCAC CANCCICGGC CICCCAAAGT GCIGGGATTA CAGGCGIGAG CAACCGCACC
TGGCCTIGAA CCCTITGAAG TATIGATGCA AAAACAAGIG GICAGCTATG GCCAAATTCG CAATTCAAAA AGATCCAAGA
AAGCAAGTIG AACATCCIGA TIGGAGATGG GACACACCCA AACGIGIGIC TIGAGGIGGC TGCAAAGTCC TCCGGICTGA
GCCAGINTAA GCAGGTITTA CCCCAGCCCA TGATTTAGAG AGATGTINAG TGCAGATCCT GAGCTCAGCA GAGAGCAACA
T

SEO ID NO: 2067: (Length of Sequence = 335 Nucleotides)

CTGGCTCTGT GGCTCAGGCT GGAATGCAGT GGGCCGAGGT TGGCTCACTG CAACCTCCAC CTCCTGATCT CAAGNCGTCC TCCCACCTCA GCCTCCAAG TAGCTGGAAC TACAGTGGAA CTACAGGTGG ACAACATCAC ACCCAGGTAA TTTTTNTNAT TTTTTTGTAGA GACGGGGTTT CACCCTGTTG CCCAGGCTGG TCTCAAACTC CTGAGCTCAA GCAATCTGCC CACCTAAGCC TCTCAAAGTG CTGGCATTAC AGGCATGAGC CACCGTGCCT GGCCTGGGAA GCTCTTTTAA CAGAGGTGAT GTAAAGTAGA AAAAGCAGTG GGCTC

SEO ID NO: 2068: (Length of Sequence = 274 Nucleotides)

GCAACCEAAT GGACACGGTA AAGAAGGAAT GGGAAGAGGC AGACCTTCAA GCTAAGAACC TCCCCAAAGC AGAGGAGCCAG ACTCTGATTC AGCACTTCCA AGCCATGGTT AAAGCTTTAG AGAAGGAAGC AGCCAGTGAG AAGCAGCAGC TNGTGGAGAC

CCACCIGGCC CGAGIGGAAG CIAIGCIGAA TGACCGCCGI CGGATGGCIC INGAGAACTA CCTGGCIGCC TTGCANTATG ACCCGCCACG GGCINATCGN ATTCINCAGG GCIT

SEO ID NO: 2069: (Length of Sequence = 321 Nucleotides)

GIGCCATCIG TITACTICIC AAATGAAAAA GAATTCAGGI CIGAGIGICC AGGAAAGGGG GIGAATIICA TAACCGCCIG
TGACAGCGAT GGGAAGGAGC CACACCCCIC CAGAGGGTAC CACCCAGCGG ACAAGIGGGG AGGAGGAAGI AGCIGGCATG
AAGCCGGCCC ACCCAACCIC CGGGAGAGAG GAAAAGGAGA ACACGGGATG AGGAGGCTIT AAATAGIATI TCATAAAAATA
AAAATGCCCA GCACICTIAG GAACCICICA TICAACCGCC TAGIITIIGI TIAAATAATI CIAATGCCAG AGCIGGGGGG
C

SEO ID NO:2070: (Length of Sequence = 161 Nucleotides)

AAAGCTGCAT AAAACAAGIT TAATTICCAA CCAGGGTCAC AGTCATCGCG TIATCCCACA TITTGAGCAA GGATAGAGAA GGTGAGTTAT TAAACATATA CAGTCTACAT TCCAGAGGAG GAACTGCAGT TACCACTATA ACACCACAGA CAAACTTTGG G

SEO ID NO:2071: (Length of Sequence = 288 Nucleotides)

GTGGAAGGC CTTCATACAT GCTTCCCATC TTCAGGAACA TCAGAGAATT CATACTGGGG AGAAACCATT CAAATGTGAT ACATGTGGTA AGAACTTCCG TCGTAGATCA GCACTTAATA ATCATTGCAT GGTCCACACA GGAGAGAAAC CATACAAATG TGAGGACTGT GGTAAGTGTT TCACTTGTAG CTCAAACCTT CGTATCCATC AAAGGGTCCA CACAGGAGA AAACCTTACA AGTGTGAAGA ATGTGGTAAG TGCTTTATTC AGCCTTCACA ATTTCAGG

SEO ID NO:2072: (Length of Sequence = 284 Nucleotides)

TCTTGTCTTC AGACCCCTTT GCCGTATTGT CCCTCCTAAC TGGGACCTAA GCTAAGACTC AAGGGCTGCT CCCATGCCCT
TCAGTATCCC CCATAAAATC TAACTACACA TTAGAAACTC AAAGAATAGC ATAGGCATGA TCCATCACCT GCAACAGAAG
CAGTGAGGAG ACTTAAGCCA GGGTTCCTINC AAGNGATTNC ACCGACCNIT CCTGCATCTC TGNATGCCGG ACTCCTAAGC
ATTTACTCAG ATTTTAAACA GCACATAATG CCATGGCGAG GATG

SEO ID NO:2073: (Length of Sequence = 270 Nucleotides)

GGAGCGATAC GCCCCTGTCG CGAAGGACCT GGCGTCTAGA GATGTGGTGT CTCCGTCCAT GACTCTGGAG ATCCGAGAAG
GAAGAGGCTG TGGCCCTGAG AAAGATCACG TCTACCTGCA GCTGCACCAC CTACCTCCAG AGCAGCTGGC CACGCGCCTG
CCTGGCATTT NANAGACAGC CATGATCTTC GCTGGCGTGG ACGTCACGAA GGAGCCGATC CCTGTCCTCC CCACCGTGCA
TTATAACATG GGCGGCATTC CCACCAACTA

SEO ID NO:2074: (Length of Sequence = 278 Mucleotides)

GCACATGCCA TCAGTCCTGG CTAATTITTG TATTTTTAGT AGAGACGGGG TTTCGCCATG TTGGCCAGGC TGGTCTCGAT CTCCTGACCT CAGCTGATCT GCCCACCTCG GCCTCCCAAA GTGCTGCGAT TATAGACAGG AGCCACCGNC CCCGACCCTC TCTCACTTCT CAAATCTCTT TCCTTTTTCC ACCITCTAGG TGTCAAAGAC AGTGGATGGT CTCTGAGGTT CAAAACCAAG CTGACCGGGT AAGTATTTAC AGCAAAGCAT CCAATGGG

SEC ID NO: 2075: (Length of Sequence = 232 Nucleonides)

GTCTCTAGGA TTCACTCAAA CCCAGGATCA CGGTTTTGTA ATGTTATCAA GGCATGATTT TGGATTTCAG AGCTGGCCCA GTGAACAACA AGCAATCAAG CATTCCTTTC TCTTTCTTTC TCTCTCTCAC ATATACACAC ACACTCTTTC TCTCTCACGT TACTITCACT GICACTITCT CICTACTGGA TAACAGGCCA AAAGTACTGG CACTCATCTT TCACTITCTT CC

SEQ ID NO: 2076: (Length of Sequence = 223 Nucleotides)

GICACGAGGT CAGGAGATCA AGACCATCCT GGCTAACACA GIGAAACCTC ATCTCIGATC TATTCAGGGC TCINACITCT TCCIGGTTTA GICITGGGIG GGIGIATGIG TCCAGAAATG TATTGATTTC TTCTAGATTT CIAGITTATT TGNGTAGAGG TGTTTATTCT CIGATGGTAG TTTTGTATTTC TATGGGATCA ACGGIGATAT GCICTITTATC ATT

SEO ID NO:2077: (Length of Sequence = 323 Nucleotides)

GTCCCCCTTC CCCTCTTGTG AGACCAGGCT CTGTCTCAGG AACAGGCCTG AGGGAGGAGG AGCCACGTTC CTCCTTCCTT
GGAGCCCTGA GGTGGCCAGG CTGTCCCCAC ATAAAAGCATG ACATCCAGGT GCCAGCTGGC TAAGAAATGG AGCCTGAGGC
TGCAGCTCAC CACCTGTACC TCACAGATGT CCATTCAGAG GAAAGAAGGG TGCTCCAAAAC GCCAGGCCCC CAAGGAGCAC
AGACTCAGGG TCCAGGCAGG TTCAGTGCTA GTAGGCAGGT GGGCACTGCT GTCCAGGAAA ACCTGGTGGG CAGCTGTTTT
CCC

SEQ ID NO:2078: (Length of Sequence = 310 Nucleotides)

AATTIGCAGT TGITAAATCA AACCTACTGA CATTIATAGT CCCTTACTTT CICTICTITC TICCATIGTA AATGICIGAA ATGICGTACA GICATACTIC CCACTGTATT TITAGGITTT ACTCTCATAC TICAATAATC ACTACCACCC TITATTICAA TAAAAGITTT AAGICAGTGC TGATTITTTIG GIAGCTCCCA TITTCTGATA TATTITGTCAT GTACATATGC AAGIGTATGT AATGTAGGTG TGCATCTATA TATACCCCACA TATACATATA TACATATACA TATATATGTC CATATACACG

SEO ID NO:2079: (Length of Sequence = 281 Nucleotides)

GAGACCTGCC AGAAGATTAA AAAAAAGAAT GAGAGAAAAG CCCAGTTAGT GGTGTGCAAA CTTACTTCCT TTAAATGTCC CATGGATGTA GGACAGTGCC ATGTTTCAAG ATGCCTGTGA GCTAGGTCTT CAAGATTTAT AGAATGTTAC TTATGAACAA AATATAATTA TTTATGGTAC AATTCTTGTA CTTTAGCAAA TCTGGAGTTA GTTCATAGTC AAAGTCAGTT AATATTTCTT AGAGGAAAGT TTTGGCTTTT TGTGGCAACA TTTTTATAGC T

SEO ID NO:2080: (Length of Sequence = 311 Nucleotides)

ATTAAAAAGA ATATTATTTA TTATCINCTT TATTAATACT CACATGIAAC CTTTGCTTTT TACACAAAAG TCTGCTTTAG
AAGAATGCCT CCNCGGCTTA TCATGCCCAA TGGGGCTTT TGTTTCTGGA CCACTTCCCC TTTCTCCACC CCCACCCCCA
CATCCAAATT ACTCTTAACA TGTTCACAGA TACCACGNAT ATTTTGTAAA CAAGNITTGG GTTACTGGAA CTTGATTTCA
TTAACATCCC ACTTCAAAAT GGAAGGCAGG TGGAGGGCAG GGTAAGGNAA TAGGGGGAAA GAGGGCAAGA G

SEQ ID NO: 2081: (Length of Sequence = 207 Nucleotides)

GGACGCACGC TCGCTGCCAT CACCGCTGGG TGGTTTTTTC CCCCTAACTT TTTACTTAGC CTTTTTGGTT TGINICCCCA CCCCCACCTC CTCACCCCCT TTCCAGTTCT TCTTCAGGCC CCTCCCAGAC GCACCCCAGC GGCCCCTGCCA GCCCCTGCCT CCAGCCTCCA GCCTCACCTT TGIGCCCAGA CTCGCATTTG GAAGACT

SEO ID NO:2082: (Length of Sequence = 260 Nucleotides)

TTAAAAGAA GTGCATACIT ATTTGCAAGG AAAACAATG GAATAGACAA AAATTTTAGA ATATAAAGAC TTTTTTNCAT
TTATGTATGT GTTTACAATT CAAAATAATA AAGCTAGTTA AAAGTCAATA CATATTAGAT ATATTCAAAT ATTTTNCCAA
ATAAATTTCG ATCTTATCAG TTAACACCCA TAGCAAAAGA CTAAGGAGTA TTTGTATAAC ATTAGGGTAT TTGACCTCAT
ATTCTATTCA TTTGGGTTTA

SEO ID NO:2083: (Length of Sequence = 257 Nucleotides)

AGITICATAT GITTATTAAA CCAAGCATGA GGCCCTTCTG TGCACAGGGC CCTGTGTGAC GGCATGGGAG GCGTGCTCAT
GAGGCTGGGC GTGCCCGCCA GAGACCTTTC TAAAATGCAG ATTCACGACT CTCCTCCTCA AGCCACCCTA GTGGCCAGTG
GGGTCATTTC GGATCAGAGA TTCCTGGAAT AGATCTAACT AAGATGGTAG ATATTATTTT AAATAATGCC TTTTTINAGGA
ACTAGCTGCT AGGCTCT

SEQ ID NO: 2084: (Length of Sequence = 255 Nucleotides)

TATTATACAG CATTGTCAAG ATTATTTGAC AAAAGGCAGT AACAAGCCGA AGGAAAACAC ATTTACAAGA AGCTGAACAA CTTGTATCAG AACATACATC AAGGTGAAGA GTTTCGGCCC TCTTGGTATA GGGTATGTAT GTGTACATCT CCAATTTTGA ACAATGATGA CATAAGGNCT AATACTCTAT TTATTCAGGN GACCCCATAA TCAGGATAAT AGTAGGCATT CAGAGTAATA AAGTGATCAC AGTTG

SEQ ID NO:2085: (Length of Sequence = 290 Nucleotides)

GGACGCACGC TCGCTGCCAT CACCGCTGGG TGGTTTTTC CCCCTAACTT TTTACTTAGC CTTTTTGGTT TGTGTCCCCA CCCCCACCTC CTCACCCCCT TTCCAGTTCT TCTTCAGGCC CCTCCCAGAC GCACCCCAGC GGCCCTGCA GCCCTGCCT CCAGCCTCCA GCCTCACCTT TGTGCCCAGA CTCGCATTTG GAAGACTCCA CCTCCCGCCC AGGCCTGGGC TGTTGGGCCGG TTGGAGATTC AGGTTTTAAT CCACACAAGC CCCAGTGAGG GGTGAAGCAT

SEO ID NO:2086: (Length of Sequence = 342 Nucleotides)

AGITTCATAT GITTATTAAA CCAAGCATGA GGCCCTTCTG TGCACAGGGC CCTGTGTGAC GGCATGGGAG GCGTGCTCAT
GAGGCTGGGC GTGCCCGCCA GAGACCTTC TAAAATGCAG ATTCACGACT CTCCTCCTCA AGCCACCCTA GTGGCCAGTG
GGGTCATTTC GGATCAGAGA TTCCTGGAAT AGATCTAACT AAGATGGTAG ATATTATTTT AAATAATGCC TTTTTGAGGA
ACTAGCTGCT AGGCTCTCTA TCCTGGGAGA AGAAGGTGAA GGTTCCGCAA TATCAATTTT CCCAACTCAG CCAAGATTTT
CCCAGCATCT NCAGGACAAG TG

SEO ID NO:2087: (Length of Sequence = 306 Nucleotides)

TATTATACAG CATTGTCAAG ATTATTTGAC AAAAGGCAGT AACAAGCCGA AGAAAACACA TITTACAAGAA GCTGAACAAC
TTGTATCAGA ACATACATCA AGGTGAAGAG TTTCGGCCCT CTTGGTATAG GGTATGTATG TGTACATCTC CAATTTTGAA
CAATGATGAC ATAAGGNCTA ATACTCTATT TATTCAGGAG ACCCCATAAT CAGGATAATA GTAGGCATTC AGAGTAATAA
AGTGATCACA GTTGAATGAA CGTGTTCACC AAAAGTCTTA GACCAACCTG ATATCATCTT ACACTT

SEO ID NO: 2088: (Length of Sequence = 326 Nucleotides)

SEC ID NO:2089: (Length of Sequence = 291 Nucleotides)

GGGTTTCCCT TTCCACTCAT CGGAGATTCA GAGGGATGAG CTGGCACCAG CTGGGACAGG GGTGTCCCGT GAGGCTGTAT CGGGTCTGCT GATCATGGGA GCGGGCGGAG GCTCCCTCAT CGTCCTCTCC ATGCTGCTCC TGCGCAGGAA GAAGCCCTAC

GGGGCTATCA GCCATGGCGT GGTGGAGGTG GACCCCATGC TGACCCTGGA GGAGCAGCAG CTCCGCGAAC TNCAGCGGCA CGGCTATGAG AACCCCACTT ACCGCTTCCT GGAGGAACGA CCCTGACCCG G

SEO ID NO:2090: (Length of Sequence = 293 Nucleotides)

TTATGIGGAA TACCACAGG CCTGGTACAT GGCTGAACTC TTCCCCTTCA TCCTGCTTGG GGTCTTCGGG GGCTTGIGGG GAACCCTCTT CATCCGCTGC AACATCGCCT GGTGCAGGAG GCGCAAGACC ACCAGGCTGG GGAAGTACCC GGTGCTGGAG GTCATTGTGG TGACTGCCAT CACTGCCATC ATTGCCTACC CCAATCCCTA CACACGCCAG AGCACCAGGG AGCTCATTTC TGAGCTGTTC AATGACTGTG GAGCCCTTGA GTCTTCCCAG CTCTGTGACT ACA

SEO ID NO:2091: (Length of Sequence = 274 Nucleotides)

CTTTIGGAAT GGTCAAACAA TTTAAGICAA ATGITTTAAT GGIGCAATTA AAATAAGGGI TCAAACATGI TITCAATATA
TTAATINCTT TAAAGICATG TTCAGGCAAG GIGCIGITTA AAAAACCACT ATTAGCITTG TCCACACATG TAAGITATCA
AAAGITACCA AGGIAATTTT GACGITGAAT GCAGCITTAA ACAATAAAAA AATGGIATTA GGITTACITC TCGAAGCAAA
GAGAGCCCCC AACCITGIAA ACTAAACATT CTGA

SEQ ID NO:2092: (Length of Sequence = 290 Nucleotides)

GETACETAGE ACECTEGCCC TYTCCTCCGG CCGENTCTGG TCAGACACAA TCATEGTCTC CACCACGAGG TGTGCAATGC CTGCNAGGGT GGTTTGCTCC AGGTCCAGGA GGGCAGATCC ATGGGCGATG GTCCTCCTGA GCTCCAGAAG GCTACGGAAG GAGAGGGAGG CAACATGGGG CTTCCCCCAG CGCTCCGTCT CCTCCTCCAC GTCCTCCTCA AACTTGATCC AGGGGGCCGT CTCCCCGCCAG TGGGGCCTCT CGCCCCAGC TCGGTCAGCT CAGCATCAGCT CAGCATCAGCT

SEQ ID NO:2093: (Length of Sequence = 323 Nucleotides)

AGCTACACTG ATACAAGTGG ACCTAAAGAA ACGAGTTCCG CTACTCCGGG ACGAGACTCC AAAACCATCC AAAACGGATC
AGAAAGTGGG CGTGGGAGGC AGAAATCTCC TGCACAGAGT GACAGCACAA CACAGAGAAG AACTGTAGGC AAAAAACAAC
CCAAAAAGGC TGAGAAGGCA GCTGCTGAAG AGCCTCGTGG AGGCCTGAAG ATAGAAAGTG AAACCCCTGT AGACTTGGCT
AGCAGCATGC CCTCCAGCAG ACACAAAGCA GCCACCAAAG GCTCAAGGAA ACCCAATATA AAGAAGGAGT CTAAGTCTTC
CCC

SEQ ID NO:2094: (Length of Sequence = 255 Nucleotides)

AAGGATGITT TGGITCCCIG CCTCAAGGCC GGCCATGIGG GAGTTGIATC TGIGGAGITC ATTGCCCCAG CCTTGGAGGG
AACGIATACT TCCCATTGGC GTCTITCICA CAAAGGCCAG CAATTTGGGC CTCGGGTCTG GTGCAGTATC ATAGTAGATC
CTTTCCCCTC CGAAGAGAGC CCTGATAACA TTGAAAAGGG CATGATCAGC TCAAGCAAAA CTGATGATCT CACCTGCCAG
CAAGAGGAAA CTTTT

SEO ID NO:2095: (Length of Sequence = 305 Mucleotides)

GCACTCCAGC CTGGGCAACA AGAGCGAAAC TCCATCTCAA AAAAACAAAG AAAGAAACTN CTGAAGTCGG GGGCTGCTAG AGGATTTTCA GGAAGGGTCA ACACAGGCCT CACTTCCAGT CCCTCATTTC CCAGCTCACA GAGTCACCAG AGGGTGAGAA GCAGAACGTG CCAGCAAAGA GGGAAAAGGC CACAGAACCA CCTTNTCCTC AATTACAAAG GGGTGCATTT CAGAGGAGGG AATAGGGATG GAGAGGAGGA ±AACACCTTC CCAGGAGCCA GATAAATTCA AAGTCACCAA GATGG

SEQ ID NO: 2096: (Length of Sequence = 327 Nucleotides)

CIAGATATAA CIACCCITCI CIATTCCICA CCIAAATCCI TATACIGCIG ATGACTITGG AAAATAGIAC AGGGITTIAC
AGICIAATCA TGACAATACA TCTCCAGGAT CCITGAGCCA AATACATTCC TCAGAATACT TITITITAAAA AACTGAAATT
GATTACTIGI ACITTGICAT CACCAAAAAT ATCIGIAGCA AGACATACIG TTCTCAGCAT CCACTTCTAC CATCCTCACT
ATTGIAACIC ACAGTAGACT ATGCCICCIA CITTACIGAA AAGATACAAA CCATTACCTA GCAATCATTC TTCCACCTTA
AACATAT

SEO ID NO:2097: (Length of Sequence = 296 Nucleotides)

CACCCTGCTG AGGTCAATTT CGTCACTGAT GCCTCGGGTC ACATAGGCCC TGATGACCCA GATTTCACAC AGAGGTCAGT
ACATCGGTCA ACTTTCCTCC CAGGAGGGGC CGGGGCTGGT GGGCCATGCC CACTCCGTGC CACATGCCTA GCATTCAGAG
CTTTGTAAGG AAGCCCTGTT CTAAATGCTC AGGTCCCACC CTTCCTTGTC AAGAGAAGCC ATGGGCTTCC TGCTCCTGTG
TCACAGTGTG CCACTTGAAG GGTGGCTCTT CCCCATTCTT CTTCCATGGG GGCCAG

SEO ID NO:2098: (Length of Sequence = 324 Nucleotides)

ATTGGITTIN TIGAGIGITI TCTTCTTTTI NITTGITTIC AACATACTTA CIGCGIATAA AGICATGCAA AGAAAACAGI GCAGACAGTA GATCCTAGIG GATGIGCCAA GGIATTCCAC TCAGAGICAA TCCCAGGGAA AGAGGGAAAG AGAAAAAGAA AGAGAGAATG CGAACCCGAG GCIGCAGGAT GAGGCATGAA GAGIAGAAAT TCCCAGIGCT TIGCIGIGGI CATCAGACGC CAAGGGGAGA GAGGCAATNA AGACACACGC TCACGGGCCC CCCAGAGGIG GGIGGGGGGT GCIGGGGGGC GGCACACAGA TATG

SEQ_ID_NO:2099: (Length of Sequence = 299 Nucleotides)

GAAACCGICA GIAAGGAGCI CITITATCTIT ACCITCCCAC TCCAAACCTA CITGCTAGCT GITCTIATCA TTGCCTCCIT
TTTCTCTGIC ACAAAAATGI GITCCATCIT AATGAACACA TITCATTAAT GICCTTCTTA ATGAAGGACA GICCCTITCC
CIGIGCTGIG AATCCCATAG TAATGACATT AGCITAAGIT TTCTGAGCAC TTGCTATCTG CCAGTTCCTC CCATGAATTA
TCTTGCTTAA GCTTTGCAGI ATACCTGIGA AATAGGTGGC AGTAGTTGTC CCACCATAC

SEO ID NO:2100: (Length of Sequence = 308 Nucleotides)

GGCAGCITAT TITIGATIGG TICACAATGI GGATCAAACA GGAAAATCIG TIATCATCAA CAAGACCAGC AGCACCAGAA
TNINCOGAGI CITICCAGCAG TICAGGCICC TICAGGCICC TICCAGCACC CICAGCAGCA CACCCCTAGT
CICAGGIGIG GCAGCIGGCI CICCAGGCIG TIGCCCTTAT CCAGAGAATG GAATAGGGGG CCAGGITGCI CCCAGCAGCA
CCAGCIACAT CCTCCTTCCA CITIGAAGCIG CAACAGGCAT CCCGCCTGGG AAGCAATCCT TICTITAAT

SEO ID NO:2101: (Length of Sequence = 291 Nucleotides)

GATGATGATT GCATGGGGTT TGATGCTACA CTGGATCATA GAGTGTGGGT TCTTTCTTAC ATGINITGGT AGATAAATGT CATAGACTGA TCCTGAATCC ACATCAACAG CATGGAATCC AGCACAGGAT CCATAGATCA CTTTCAACCT CTGGCCTTCC TCAACAGTGA GATCCACCAG TAATGGCTTA TGTACCAATT CTCCAAATGA CTTAAAGGCC ATAAATTTGG TGATATGGCT TTGGAGCCCCA CGCATAGGAC TTCCACAGAA CTTTTTCAAA GGCAATCACC C

SEO ID NO:2102: (Length of Sequence = 323 Nucleotides)

GATGATGATT GUATGGGGT TGATGCTACA CTGGATCATA GAGTGTGGGT TCTTTCTTAC ATGTGTTGGT AGATAAATGT CATAGACTGA TCCTGAATCC ACATCAACAG CATGGAATCC ACCACAGTAT CCATAGATCA CTTVAACCT CTGGCCTTCC TCAACAGTGA GATCCACCAG TAATGGCTTA TGTACCAATT CTCCAAATGA CTTAAAGGCC ATAAATTTGT GATATGGCTT

444

TGGTGCCCAC GCATAGACTT CCACAGAACT CTTCAAAGCA ATCACCAGAA ATTTGATTCT TTCATATTTT ACAACTTTAT

SEQ ID NO:2103: (Length of Sequence = 270 Nucleotides)

CCTTTCACTC CCCCGCCCTG GGCCTCTGCT CTCTTGCCTG GNITCCTTCT TITTTGAGGG AAAGAGGGTG GGGCTGCAGG CAGTCTACTG GCAGGACGGG AGGCTGAGTC CTCAGGGTCT CACACCCTCA GTGCTGATGC CATGCCAACT GCCTGGGACA ACACCAACAC GTAAGGACCT AATTAAACCA AACCAGAGTC GGGTGTAGAC CAGCCCTGGG ATTTCCAGCT NTGACTNGGC CAGGGCACAC GTTGGTCTCG GCAGTGGCTG

SEO ID NO:2104: (Length of Sequence = 367 Nucleotides)

CCTITICACTO COCCECCTE GECCTCTECT CTCTTECCTE GCTTCCTTCT TTTTTGAGGG AAAGAGGGTG GGGCTGCAGG
CAGTCTACTG GCAGGACGGG AGGCTGAGTC CTCAGGGTCT CACACCCTCA GTGCTGATGC CATGCCAACT GCCTGGGACA
ACACCAACAC GTAAGGACCT AATTAAACCA AACCAGAGTC GGGTGTAGAC CAGCCCTGGG ATTTCCAGCT GTGACTGGGC
CAGGGCACAC GTTGGTCTCG GCAGTGGCTG TAAGGTCACC TTCCTTNCTC TGGATGCTGG TTTCAACCAT CTATATATGG
CATCCACGCA TGGGATCTGC AAGCTGGAGC CCTCCTACCC GCAGCTT

SEO ID NO:2105: (Length of Sequence = 288 Nucleotides)

GCAAAATTAC TGAAACTACT ACTITGGGCT CAGAACGAGC TGGACCAGAA GAAAGTAAAA TATCCCAAAA TGACAGACCT CAGCAAGGGT GTGATTGAGG AGCCCAAGTA GCGCCTGCNC TTGCNTGGTG GATCCAACAC CAGCCCTGCG TCGTGGGACT TGCCTCANAT CAGCCTGCGA CTGCAAGATT CITACTGCAG TAGAGAACTC TTTTTCTCCC TTGTACTTTT TTTTGACCTG GNATCTTTTT ATAGGGAAAA ATGGCCTTTG TAGGCAGTGG AAAACTTG

SEO ID NO:2106: (Length of Sequence = 349 Nucleotides)

GCAAAATTAC TGAAACTACT ACTITGGGCT CAGAACGAGC TGGACCAGAA GAAAGTAAAA TATCCCAAAA TGACAGACCT CAGCAAAGGGT GTGATTGAGG AGCCCAAGTA GCGCCTGCNC TTGCNTGGTG GATCCAACAC CAGCCCTGCG TCGTGGGACT TGCCTCAGAT CAGCCTGCGA CTGCAAGAAT CTTACTGCAG TAGAGAAACTC TTTTTCTCCC TTGTACTTTT TTTTGACCTG GCATCTTTTT ATAGGGAAAA ATGGCCTTTG TAGGCAGTGG AAAACTTGCA AGGAAAGCTG CCGTCTCTTT TGGCAGTCTT GATGCAGAGC CTGCACTCTG GCACTCGCT

SEO ID NO:2107: (Length of Sequence = 329 Nucleotides)

GTGACAAGCT CCAGAAGCCC GCNTCGCAAC ANCCAGGAGG GCCAGGCCAC TCCAGGCAGG AGGCAGTGGG CTGGCAGCCA
CCCTGGGCAC AGAAGAGCAG ACGCAGACAG TGCTGGGCAA CGAGGGGGTT TNTTCATGGG CCCGCCTGCC CTGTCCCTCC
CCCCAGGTCC CCACCTTCTA GGGTTAAAGT GCAGCTGGGA GGGAGGAGC AGGCAGAATT NGGGAGCTAG AGAGAGCCCA
AGTGAACCCT GACTGTCCAC GCAAGTCCCA TGTCCTCCTC GTCCTGGAGT TCCTCGAGGT TCAGCCAGCC CATCCCGCCT
AGGGCCTCT

SEO ID NO:2108: (Length of Sequence = 261 Nucleotides)

THITCATGGC AGCCTGAGCA GACTAAGACA GCAGCTAACA CAGCAAGATC ATACCAGTTA ACCITCCTGG TTAGAAGACC TGAGCCTCCT GACTTCCGGT CACTGGATAC TCTCTGTNAG GCTCATGATT TAAACTCTAT AGTCACTGCT GGCTTGGAAA CCTCTAACTC TCTCTGCCTC TTGACAGTGT TCCCTCAAGG GAGTCCATTA GCCAGGACTA GGTTACATGC CCCTGTGTTA GCTGTGAGGG ACAAGGCAGA G

SEQ ID NO:2109: (Length of Sequence = 329 Nucleotides)

TTITCATGGC AGCCTGAGCA GACTAAGACA GCAGCTAACA CAGCAAGATC ATACCAGTTA ACCTTCCTGG TTAGAAGACC
TGAGCCTCCT GACTTCCGGT CACTGGATAC TCTCTGTTAG GCTCATGATT TAAACTCTGT AGTCACTGCT GGCTTGGAAA
CCTCTAACTC TCTCTGCCTC TTGACAGTGT TCCCTCAAGG GAGTCCATTA GCCAGGACTA GGTTACATGC CCCTGTGTTA
GCTGTGAGGG ACAAGGCAGA GAAATAACTG CCCAAGTTCA GCTTCCCATA ATGTTTGGGG GATGCTATGA CTCAACTTTG
ATCTATTTT

SEO ID NO:2110: (Length of Sequence = 271 Nucleotides)

SEO ID NO:2111: (Length of Sequence = 315 Nucleotides)

GGCTTGAGCA GACAGAACGG GGAAGACTCC ACTCTGTCCC GAGGGGCCAG CCGCACGTNC NCCCAGGGCC ACCCTGCCCT GAGGTCCTTG TGTGGCCGCC CTGGCTTGGC AGCCCTGCCCC ACGCTGCCCC CGCAAACAAT GGTGTGTGCG TTTTTACAGC CCTTTTTAGG AACCCAATAT GGGCATAAAT GTAACACCTG TAGCGGGGC AGATTCTCTG TATGTNCAGT TAACAAATTA TTTGTAATGT ATTTTTTTAG AAATCTTAAA ATTGCCTTTG CACTGAAGTA TTTTCATAGC TGTTTATATC TCTTT

SEO ID NO:2112: (Length of Sequence = 275 Nucleotides)

GCAAGANAGA CCAAAACCTA ACCTGAGTTA CAAGAAACAA GACAGTAATG GCTATAAAGG GAGTGACCAG GAGCAACTGG
GACACTCCTT TACCTCCCAT ATCCAATGTA TGINTTTCAC AGAAAAACAA CAAAATTAAC AAATTCACAA AATACAACAG
CTAGAATTAC AAAATCCATT CATCCAAGGG TGGTAGAAGG CAGGATGGNA AGGTGGAAGG GTAAATNGCA CAGGGAGAAA
AACAAAGTGT TCCAATCAGT CCAGGCACAG GGACT

SEO ID NO:2113: (Length of Sequence = 227 Nucleotides)

GECGCATCAG TGGGGGGTGC TGTCAAAATT AGTGAAATCA GATACAGTTG ATGGGCAGGG AGGGTGGGGT AAGAGACAAC
TCCAGTGCAG TGCCAGGTGG GCAGGCTCCC ACTGTTCACT TGAGACGCTC CTCCCCACTC AGGTGGGGAC AGGGACACA
CTCGCAGGGC AGGGCATTCT GGAGGTGTGG GTACAGGTGA GGGGAAATGG GAGGCACACC CAGGAGT

SEO ID NO:2114: (Length of Sequence = 339 Nucleotides)

GGCGCATCAG TGGGGGGTGC TGTCAAAATT AGTGAAATCA GATACAGTTG ATGGGCAGGG AGGGTGGGGT AAGAGACAAC
TCCAGTGCAG TGCCAGGTGG GCAGGCTCCC ACTGTTCACT TGAGACGCTC CTCCCCACTC AGGTGGGGAC AGGGGACACA
CTCGCAGGGC AGGGCATTCT GGAGGTGTGG GTACAGGTGA GGGGAAATGG GAGGCACAGC CAGGAGTGGG GCAGGAGGGA
AGGCCAGTTC GTNGGCAGGC TGAGGAGGGA ATATNACCCC CCTCAAGTCC CCAAAGTGGC AGGCAAGTTA AGGGGCCCTG
GATGAGGTGG CCCCTCATG

SEO ID NO:2115: (Length of Sequence = 262 Nucleotides)

TEGRACACAA AATTCCCTGT MITAACATTG TACATTCGGG GCCTAGCTGC CCTTGAGGAT GTCCTAGTTA CACCCTCTCT
GATACCTGTG GAGTTTAAGC ACCATTCCTA CCCCTGTGTC CCTTTTGGAGG GCGTGCAGTG GAAGCTCTTA AAGGGGATC
CTTGCTCTGC CTCTGTGGCT TTTTGTTTGG GAAAGGGAGT TMGGATTMGA GGATTTAGAT TIMAGGTCAT GATGTCAGAG
CACACCAGGA ACTCCCAAGG CT

SEO ID NO:2116: (Length of Sequence = 153 Nucleotides)

AAGAAGCGAA GAGGATTCGT GAGCTGGAGC AGCGCAANAC ACGGTGCTGG TGACAGAACT CAAAGCCAAG CTCCATGAGG AGAAGATGAA GGAGCTGCAG GCTGTGAGGG AGAACCTTAT CAAGCAGCAC GNGCAGGAAA TGTCAAGGAC GGT

SEO ID NO:2117: (Length of Sequence = 231 Nucleotides)

GAATATAATG TGTATCTINCA AGGINTOGATC CACCCTINCC CATCCTINTGG AGCTCAGAGA TTCTTGGGAG CTGAAGGTCT
TCTTAATGTC AGATCAGCAA CCCCAATCTC AGGCAGCTCG GATTCGCTGC TCTCGATCTIN CCGCTGGCCA ATGTAAAACC
AGACGCAGGC GACCCAGTGC GCGACAGGGC GAACACGGCC ATGAGCAGTG TCAGCACCAC GGCGCTGTAC T

SEQ ID NO:2118: (Length of Sequence = 309 Nucleotides)

SEQ ID NO:2119: (Length of Sequence = 308 Nucleotides)

SEO ID NO:2120: (Length of Sequence = 237 Nucleotides)

CCGCTCTCCT GACGGGAGCC CACTAGGGGG TCCTCTTTCA TCTTTGGTGT GGCCTTACCT CCCACCAAAG AGATCCGAGG CTTACTCTTC TCTCTCTGGG ACCAGCATGA CCCAGGAGTC CTTCCAGGAG AGCTCTGTGA AGGAGCTGAG GCGCCTGGAG GACCAGCTGG CCGGCCTGCA GCAGGAGCTG GCGGCTCTGG CACTGAAGCA GAGCTCGGTG GCGGAAGAAG TGGGCCT

SEQ ID NO:2121: (Length of Sequence = 224 Nucleotides)

GCGGTCAGAG GCTGAGGCCA GAGAGGTAGC AGCGGAACTN ACAGGGAGGC CAGGGGCAGA GCTGACCCTG GAGAGGGATC CTNATGTCCT AGACACATGG TTTTNITCTG CCCTGTTCCC CTTTTNTGCC CTGGGCTGGC CCCAAGAGAC CCCAGACCTT GCTCGTTTCT ACCCCCTGIN ANTITTGGAA ACGGGCAGCG ACCTTCTGCT GTTCTGGGTG GGCC

SEQ ID NO:2122: (Length of Sequence = 202 Nucleotides)

CASCIGCAGC TICCAACCAA GAAAACCICA AAGCATTAGG GAAGGAGCAG GIGIGGGGT GGGGIGGGGA GAATCCCCTA
AGCICCAGGG CCCAGGGICI AACCIGAGAG GICGGGGCTG CAGGAAGCTG GGGGAGGCTC CCGGGGCTGG GGGAAGAGAG
GCCTGCCCCC AGCAGAAACA GCAGGICTCA GCGCTACAT GI

SEO ID NO:2123: (Length of Sequence = 359 Nucleotides)

ATTCTCCTCT GTTCTCTTGA TGTGTAGGGA AATTTUAAGA ATGACTCTGA TAAAAATCTA AAAGAGAAAC ATCGAATCCT AACTGGCTGT GTGACCCTAA AACCTTACTC CGTCTCTTTG AACCTCAGAT TTCTCAGGGC TTGGCACATA GCAAGCATTT CATACTCAGA AGCTGGTACT ATTACTGTTG TGTTTTGTGG GGGGAGGTTT GTTTGTTTTG TTTGGAGACA GGATCTGGCT

TTGTTGCCCT GGCTGGAGTG AAGTGGCGCC ATCATAGCTC ACTGCAGCNT CGCCCTCCTG GGCTCCAGCG ATCCTCCCGC CTCAGTCTCC CGAGTAGCTG GGACCACCTG CGGATGCCG

SEQ ID NO:2124: (Length of Sequence = 233 Nucleotides)

GAAACGCCGT GCATCTCTTG TCTGTTGGCA GCGAGCACAT CGTNTGGAGA CACGAGTTTC TAAGCAGCTG GCACGAGGGC TGCTGACGGC ATGGGTCGTG CTTCAGGGTG GCAATACCTC TTAGGAACTT AGGGCAGGAA GCAATACTTC AGCATTGAAT GTGTGTAAAT AGTTGCTTTG AGTTGCAATT GCTATTTNCT TCTCAGTCCC AGCTCAGATC GAATTATATA TCC

SEQ ID NO:2125: (Length of Sequence = 241 Nucleotides)

GCCATGGCTT TTGGTCAGGT TCAGGGGGGC TGAGGGGGTG CTCCTCCCCT CCCCCAGGC ACTGACACAT TGAAAGGAAG CAGAGCAACA ATGACACAGC ACGGATGTGG GAAAGGGGAT CCCCCAAGGG GGCAGGATGG TCCATCTCAC CGGGGTCTCA CCAGGACTCC CCGCTCCCAC CCAGGGCCAG CACGAGCACC TCCCGTTTTC TCCCCAGTGC AGAGCGTGGG GTGACAGGAG T

SEO ID NO:2126: (Length of Sequence = 275 Nucleotides)

GTGTGCCCTC TTGCTGTGC TTACTTCATA AGGAGITGTA TCTTCCCACC TGCATTTCAA TACTGCCGGT TAGGACCTAA GTAGAAGAGC AGTAAAGGCT GATTGACCACA CAGGGGGATG GAGTTGGTCC TTGTCCATTC TCTCACCCTT GCTGTGCATG TATCAATCCT TATCCCAGAA GGTACTATTT AGACTGTATA GACTGATTTA GATTACATAC TTTAGAGGAT TAAGGAAACC ATAGAGTTTG GGCCTTGGAA CTGTTACTGC CTTGT

SEO ID NO:2127: (Length of Sequence = 296 Mucleotides)

TTCAGCCTTA TCGAAACACA TGAAGCAAAA CCATTGAAAC TGTATGTGTA CAACACAGAC ACTGATAACT GTCGAGAAGT
GATTATTACA CCAAATTCTG CATGGGGGG AGAAGGCAGC CTAGGATGTG GCATTGGATA TGGTTATTTG CATCGAATAC
CTACACGCCC ATTTGAGGAA GGAAAGAAAA TTINTCTTCC AGGACAAATG GCTGGTACAC CTATTACACC TCTTAAAGAT
GGGTTTACAG AGGTCCAGCT GTCCTCAGTT AATCCCCCGT CTTTGTCACC ACCAGG

SEQ ID NO:2128: (Length of Sequence = 322 Nucleotides)

GCATGGGAGG GAGGAAGAGA GGITTGGGGT GCGGTGGCAG GTGATATAGG GAAAGGGCTC ACGITTCAGA ATCTGTGAAC
AATTCCATTT TTCATCAGAT AGCAGAACAA CTACAACAGC AAAACCTAGA ACATCTCAGA CAGCAGCTCT TGGAGCAGCA
ACAGCCTCAA AAGGCCACTC CTCAGGATAG TCAGGAAGGA ACCTTTGGGT CAGAGCATTC AGCGTCACCA TCACAAGGGA
GTAGTCAGCA GCATTTTCTT GAACCTGAAG TCAATTTGGG ATGATTCCAT AGATATTCAG CAACAGGATA TGGATATAGG
AT

SEQ ID NO:2129: (Length of Sequence = 222 Nucleotides)

TTTAGTGGGT CTGGGGTGGG CGCGGCCCCC GGCTAACGG GCGGGTCTCC TCCTCTAGGC GCAGGAGTGC GCGGTGCTCT CCAGGCCTCC CCGGCTAGGT GGAGCGTGAC ACCGCAAAGC ACACCGTCCT ACCGAGGCGG GCCCAAGGC GCACCAGCCC CTCCCCAGAT GGAAGTGCCC GGCAGACAGC TGCCCAAGAC CTCACAGAAC AAAGATGGAC AT

SEO ID NO:2130: (Length of Sequence = 191 Nucleotides)

GREGATICT TRATTICACT GREGOGGEA GGEAACCTGG ACALGGEGGG GCACGLGGGG TGGEGAGAC CTAGCCCAGC GGGGACTAGG CAGGGGAGAG GCTGCCCCCA GGCCTGTTGA GGAGAAACTN AGGCCAGCCC TGGCGGAGAC CTAGCCCAGC GGGGTAAGGA GGGTGGGGGA AAACTGGGTC T

A ...

SEO ID NO:2138: (Length of Sequence = 305 Nucleotides)

ATGCTGAGTC GCAGTTCCGA TGTTCTTATG CTTCCATCAG CAAATCTCAA TTTGTCAAGA TTCATGACAG ATTCTTCCCC
AGCGTTTGGT TTAATTGGAG GGACTTTATC TCCAGGCCTG CATGACTCTT CGATGCTCAG GGCACATGCC CGACCAAAGA
CAACCAGGTC CAAGAGCGAG TTTNCCCCGA GGCGGTTGGC ACCATGTACC GAGGCACAGG CGGCCTCCCC ACAGGCGTAC
AGGCCGGCCA CAATCTGATC CTGGCCATTC ACGTGCCTCA GGACCTGCCC CTTGTAGTTG GTGGG

SEO ID NO:2139: (Length of Sequence = 263 Nucleotides)

CGGCCCCCAG CAACAGCTTC AGCCCTCTCC ACCAAGCTTG CCATCAAGGA GATTGCACCA CTGCACTCCA GCCTGGGTGA CAGAGTTAAG ACTCTCATGG GGACACTACT GTTCAAAAGG CCCTGGCCAA ATAACTCCCA AATGAAACAC TCAACCCAAG GATGTTTTCA GCCCACTGTT AGTGAAGCTG GGTGCAGAAT GCAAAGCCTC TAAAAGGAGA GGATACAAAG TCAGGTGAGT AGGGGCCATT GGCAATGCTC AGA

SEO ID NO:2140: (Length of Sequence = 255 Nucleotides)

CTGCTTCANT CTGCGCCCCT CAGCTGTGGC TTCCCGGCAT GCCCTGTGAC CCCAAGCCGC AGGTACAGGA AAGAAGTTTG
TGCTGGGGGA CTCAAAGACC CAGAGGTTAA TTAACAGGAA CCAGGGCCAG GGGCCTTCAT CTAGAGGTCA GTGGAGTCTC
CAGGGCACTC ATCACTGTGG CTGGGAGACT ACAGTGTCTC GGCTGCGGAC TTGTGGAAGA AGAGGGGGAA GGATGGGAGA
AGGGGTGACT GGATG

SEO ID NO:2141: (Length of Sequence = 355 Nucleotides)

TITAATTAAA TACCACTICA TAATGITATI TGCACCIAGI ACTITITITI TITIAAATAA GACATGCCAT AAGICGIGAA GITAACAAAA TATAAGCATC CGCACAGAAT ATATICTAAG GIGACITCAT TIACACCGCI TCTCAGAGAA ACACACAAGI AACCTITIGI CIGCCIATCA GCCAGIGITG AAACAGCTIT GGAATICACA TGGAAGGCTG CCGGGCTGGI TCCCCAACAC TINGCCTGATG GAGTCCTGTA TCCGNACCGI GCCGTCAAAC TGGCTGGTTT CCACTAGAAA AGCAATGGAG AGTCAGCTCT CCCTTCTTTA CCCAGCGTTC AACTCCACAC TGCAA

SEO ID NO:2142: (Length of Sequence = 391 Nucleotides)

CTGCTAAGTG CCATGAGACC TTAGCAGAGG CTGTGGGTGC CCCGCCCCAT TCCCTCCACT CACTCTTCCT TGCAGGTCGA
CCTGCCCTTC TTTGCTGAGG CCTTTCTCTG CCTCCAGAGC CTGCTTGGTC CTCAGGCTGT AAGTGCAGGC AGAGCTAATG
TCTCTCCATA GCTGCCCTCC ACCAGCCTGC TCCTGAGACA CCTGCTGGCC AGCAGCCTGA AGCAGAATCC TTTACTCAGA
TTCAGCCGCA CAGATGCTCA CTGCAGAGAT CTCCAAGGNC TGTGGTCATC CTTGAGCCCA TCTCAGATTT GTGTGGATAG
GGTGTTAGAG AACATGGAAT CAGCTGGATA GAGTGGTTCA TGCTTGTAAT NCCAGCACTT TTGGGAGGCT T

SEO ID NO:2143: (Length of Sequence = 326 Nucleotides)

GATGCAGAAC AGCITCITGC AGAAGCACCI GCTCCGGCAI CCAGCGCTGC CTGGAGGCAG GAAGGAGAG CAGGCCAGGA CACGCTGGTC TGAGATGAGG GGGAGCCCCA CGGGCCCCAG GCAGGCTAGA GGAGGCACAG GCCCTGCCAC GGCCAACTCA GGTCAGCCAG CCTGAGGCTG TGGCCTCCAA AGGGTCTGGG CGCACCCCCC AGGTCGCAGG TNINTGAGGC CAGCCAACCT GCAGAGCACT CGCGGCGTGG GTGGGCTGAG TGGAGGTGCC TGGGAAGCTG CCTAAATTCA GAAGCCTCCA CTTGCCATCG AGACTG

SEC ID NO:2144: (Length of Sequence = 357 Musleotides)

GCACCGGGCC CCAGGAGCCC ATCAGTGACA GAGTGCTCCA TGATGATGTC CTCCACCGG GTGATGTACA GCAGCGTCAN
AGCACCCCCA GGAACTGGGA NAGCAGGATG CCCAGGAGGA TGCCCGCCAT GATGGTGTAG TTGTCCATGA ACCAGATGAT

CACGGCGTTG GTGCAGCCCC GCACGTAGAT GACATCCTGC ACACTGAAAC GCTCCTTGTC GATAGTTTTN TAGCCACACA TGGTGTTGAC AACTTCTGTC GTGTTCCTGA TGCAGCAGGT GTAGGGCACC CCACAGGCCA GGGGTCCAGG GGCACTGCAG TCGTGGTACT GATTCTTGCT CCAATCTCGG TAGTCCT

SEO ID NO:2145: (Length of Sequence = 420 Nucleotides)

CGCCAGGAGC TECTAGCCAA AGCATTGGAG ACCCTACTGC TGAATGGAGT GCTAACCCTG GTGCTAGAGG AGGATGGAAC
TGCAGTGGAC AGTGAGGACT TCTTCCAGCT GCTGGAGGAT GACACGTGCC TGATGGTGTT GCAGTCTGGT CAGAGCTGGA
GCCCTACAAG GAGTGGAGTG CTGTCATATG GCCTGGGACG GGAGAGGCCC AAGCACAGCA AGGACATCGC CCGATTCAAC
TTTGACGTGT ACAAGCAAAA CCCTCGAGAC CTCTTTGGCA GCCTGAATGT CAAAGCCACA TTCTACGGGC TCTACTCTAT
GAGTTGTGAC TTTCAAGGAC TTTGGCCCAA AGAAAGTACT CAGGGAGCTC CTTCGTTTGG ACCTCCACAC TTCTGCAAGG
CCTGGGCCAT ATGTTGCTGG

SEQ ID NO:2146: (Length of Sequence = 390 Nucleotides)

CCCAAATACT GTTTCCCAAA CTATGTCGGG CGGCCGAAGC ACATGCGGGT NATGGCTGGA GCCCTGGAGG GGGACCTCTT
CATCGGACCA AAAGCAGAGG AGCACCGGGG GCTGCTGACC ATCCGCTACC CCATGGAGCA CGGCGTGGTG CGAGACTNGA
ACGACATGGA ACGCATCTNG CAGTACGTCT ACTCCAAGGT TCAGCTGCAG ACCTTCTCGG AGGAGCATCC TGTGCTCCTC
ACGGAGGCCC CGCTCAACCC GAGTAAGAAC CGGGAGAAGG CGGCAGAGGT GTTCTTTGAG ACCTTCAACG TGCCGGCCCT
GTTCATCTCC ATGCAGGCTG TTCTCAGTCT GTACGNAACA GGACGCACGA CAGGAGTGGT TCTAGACTCA

SEO ID NO:2147: (Length of Sequence = 219 Nucleotides)

TITGIGGIIG GAGAGAACT GGIGIICTGC CCGCCTCTGC TIGGICACAG ACAGCTCCAG CAAGAGCAGI TGITAAAAGI GCCAAGCGIG TGIATCACTG TGACAAGCCG TITGCTTACT GCCCTGTTCC CTINCAGCCA AACCAGCTGA TGAAGAACTG CTGCCAGGNG GGICCTACAG CAGGICACAA ATGACCTAGI TTCATTITAA GCAGACAGA

SEQ ID NO:2148: (Length of Sequence = 353 Nucleotides)

GAAATCTITA TIACAAAAAT ATTITGCAAG CCAAAAAGIT TAAGITGCAA CTATATACAA AATGGGGCCT GTITCCTTCC CAGCAGTCTT AAAATAAACT CCTGAAACCA TGCTCCTTCC GCAGGTTGGT TCGACCTCTT CCTTTTCCTG GGGTTCAATA CACAAGGTAT GTGGATTCTC CAGGTTGCCA GGCTAAAGCT AAAGCTATAC ATCTTCCTTG GCCTTATTCC CTTATTTCCC CCTCCAAGAA TTAAAAAAATA AAATAAAATG AAAATGGCAC CAAGAAAACA TTCTTTTAAA ATACTGAATG TGTGTGTGCA TGCGTGTGCA CAGTATGTCC CTGTTCTCTG GGT

SEO ID NO:2149: (Length of Sequence = 394 Nucleotides)

GGGGAGACIT TGGGCTTINN TCATGACTGT TTGGGTCGAA GGTAGCTCAA GIGIGIGIGI GIGIGIGI GIGIGIGIGI GIGIGIGI GIGIGIGI GIGIGIGIGI GIGIGIGIGI GIGIGIGIGI GIGIGIGIGI GIGIGIGI GIGIGIGI GIGIGIGI GIGIGIGIGI GIGIGIGI GIGIGIGI GIGIGIGI GIGIGIGI GIGIGIGI GIGIGIGI GIGIGIGI GIGIGIGI GIGIGI GIGIGI GIGIGIGI GIGIGIGI GIGIGIGI GIGIGIGI GIGIGIGI GIGIGIGI GIGIGIGI G

SEQ ID NO:2150: (Length of Sequence = 200 Nucleotides)

ACCTCCCTGG CCTCGGAGA CGCTGACAGC TGGGACGACA GCAGCTCCGT CAGCAGCGGC ATCAGCGACA CCLTAGACAA CCTCAGCACT GATGACATCA ACACCAGCTC CTCCATCAGC TCTTATGCCA ACACACCTGC CTCCTCTCGA AAAAACCTGG ATGTGCAGAC TGATGCTGAG AAGCACTCAC AGGTGGAGAG SEO ID NO:2151: (Length of Sequence = 369 Nucleotides)

GIGGGCCCCA GICTICIGA AACCIGINAT CACACITCGG GCACIGICCC CICTACAGIC AATCIGIGIT ITCAGAAGIG
GCCCCAGGIT CACTCGICIT ACAGCAGICC TAAAGAGCCG GCIGCCCITI CCCTAGGCIT CCTIGCICIT NAGGGCTAAA
TICCAGCCCT CCTACCCCAG TGCCACITGG GIAAAAATAC TCIGCICCIC TCACGIFIGC TAATAAGCCC GGGCTCCGAC
TACCACCGIT CGGGGAAGG GAGCCCCITA CCGICATTGC TGGGICCGCT CCGGGAAAAC ATGIGCCGGA CCTGACITGT
GCGGCGGCAT CITTCCGGAA ATGCCGITTT TGITTCCITC TAAGGGIGT

SEO ID NO:2152: (Length of Sequence = 312 Nucleotides)

TTCACAAACA AATTGTGGGA GAAACACACC TTCCCAGCAA TAGAAAATCT CTATAAAGTG CATTTTGCCT GCAACCATCT CTCCCCATG CTGGCCCTTG GGTCAGGATT TGAGGCACTG TTCCGAGGGA GCCCTCAGGG CCACCTGAGC TGGGAGAAGG GAGGCATGAA GCCACCATGG AGCTCCAGGC TACTGGACAT ACCCTCTCTA CCCTGCCCTT CCCTNTTGGC TCCAGGAGTG CACTGCCTGA CTCCACTGGC AGGTTGATCT GGGAACGGCC TNGGCATGCT AGGGATGGTG GAGAAGTAGG CG

SEO ID NO:2153: (Length of Sequence = 325 Mucleotides)

CCCAGACCCA GAATGTAAAT NAGGCCAAAA TGGCCACTTC CCAGGCTGAC ATAGAGACCG ACCCAGGTAT CTNTGAACCT
GACGGTGCAA CTGCACAGAC ATCAGCAGAT GGTTCCCAGG CTCAGAATCT GGAGTCCCGG ACAATAATTC GGGGCAAGAG
GACCCGCAAG ATTAATAACT TGAATGTTGA AGAGAACAGC AGTGGGGGAT CAGAGGCGGG CCCCACTGGC TTGCAGGGAC
CTGGNGGTCT GCACCAGTTC CAGTGACCAC TTCAGAACCC ACCTNGGNGC ACCCCCCAAT GTGCTCTGGC AGACGGCATT
GGCTT

SEO ID NO:2154: (Length of Sequence = 326 Nucleotides)

ATCATTTAAT TAACATCTTT AAATGAAACA CAGTITICIT CATGIGICIC ACICAGGCIT CAGGGCAGAG GGAATGGATT
TITAGACATA TCAAAGACTC AAAAATITAA AGAAATATAT ATATGIATAT ATATACITICT AACATITIAT GGAAATTAAA
AATCAGAGGC TITIGGICIC TCCATITACT CIAGGICAAG CICATITACC CCAGAGGACA AAGAAGGGCT GCCICITCIA
GACCCICCCT TCTCCTITGI CCINIGICCC ACCCAGCAGG GAAACAAGCT CAGAAGGATC CTAACAGGAT AGAGITITCCA
GTAAAT

SEO ID NO:2155: (Length of Sequence = 317 Nucleotides)

TGGATGAGGA GACCCTGAAC ACACCCTGCT ACTGNCAGCT GGAGCCCAGG GCCTGINACA TCCTGCTGGA CCAGCTGGGC
ACCTACGTTT TCACGGGCGA GTCCTATTCC CGCTCAGCAG TCAAGCGGCT CCAGCTGGCC GINTTCGCCC CCGCCTCTG
CACCTCCCTG GAGTACAGCC TCCGGGTCTA CTGCCTGGAG GACACGCCTG TAGCACTGAA GGAGGTGCTG GAGCTGGAGC
GGACTCTGGG CGGATACTTG GTGGAGGAGC CGAAACCGCT AATGTTCAAG GACAGTTACC ACAACCTTGC GGGCTCT

SEO ID NO:2156: (Length of Sequence = 372 Nucleotides)

CTTCCAGCTG GCAGCCAGT GGCCCACCCA TGTCAAGCAC TTTCCAGTGG GACTCTTCAG TGGCAGCAAG GCCACCTGAG
GCCCTGINTC CCAGCCACTT TCCCTCCTGG CACTGCCACC AGCCTCACCG AGTGGCGCGA TCTCGGCTCA CTGCAGCCTC
TGCCTCCCGG GTTCAAGCAA TTMTCCTGCC TCAGCCTCCT GAGTAGCTGG GACTATAGCC GCGTGCCGCC ATGCCCAGCT
AATTTTTGTA TTTTTAGTAG AGACAGGATT TAACTATGTT GGCCAGGCTG GTCTTGATTT CCTGACCTCG TGATCCGTNC
TCCTCAGGCT TCCAAAAATG CTCGGATTAT AGGCATGAGC CACCACAACC GG

SEO ID NO:2157: (Length of Sequence = 351 Nucleotides)

CTGGCTAACA TGGTGAAATC CCGTCTCTAC TAAAAGTACA AAAAATTAGC TGGGCGTGGT GGTGGGCACC TGTAGCCCCA
GCTACTTGGG AGGCTGAGGC AGGAGAATGG CGTGAGGCAA CAGTGCAGCC TGGGCAACAG TGCACCTCCT CCATCTCTAC
CAGCGTCCCC TCCAGTCTGC ACGGGGCAGT CCTCCTGGGC TTGACCTCTC TGTACCCACA GCTGGGGGCC AGGCAGCCCC
CCTCTATCCC TCCCAGCACC TACTACATCG NCCINCACAT CCCTGATTCC TGTTGTTATG GAAACTNTTG CCAGAGATGG
AGGTTCTCTC GGAGTATCTG GGAACTGTGC C

SEO ID NO:2158: (Length of Sequence = 280 Nucleotides)

CAGCTCCTGA GGACCGCTGC AGTGATGACA CAGGACTATT GCATCAGCAT CGTGCTCACA GGGAATCAGA GCTCAGCCAG
GAGAGGTCCA AGAATGACAG AACCATGAGC ACTCCTACCA AAACTCAGCT CTGCTCAGCC AAATCAACAA TTCAACCCAA
CAGGNCAACT CCTAACACAT CCCATCCAGA CAGACATTAG AGGCGCACAG CAGATGAACC TCCTACTTAC ACTGTCCAAG
GAAGCTGGAC TATCAATTCC CAGTAAAAGT GGGGGAAAGG

SEQ ID NO:2159: (Length of Sequence = 342 Nucleotides)

CTTGTGCGTT TCTCCTACCA GATTGTGCAT GCCTCCTGTG GGCAGAGCCT GTNCTGACTT GCTCCTGGGT CTCCAGCATC

ACCCAGTCTG GAGCTGAGGA CCTGGGTACC TACAGATTTC CTTCCACACT GTCAGAATTG AGATGAAGGA AGCCCAGAGA

AATCAAGTAC CCTCCACCAG GCAGAGCAAA GTCCTGGGTG CCCAAAATCC AGGGAAGGCA AGGGCTGGGG GTACAAGCAG

AGGATCTGAA GAGGTATATG AGAGTNGCCA GCACAGACCT GGCATAAGCT TGGTGCTCAG TGAAGGTTAC CTGATGTTGC

TGGGCACCAG GGGTGATGCA GT

SEQ ID NO:2160: (Length of Sequence = 376 Nucleotides)

SEO ID NO:2161: (Length of Sequence = 404 Nucleotides)

CCITCCITCG GITTCAACTG GACITCIATC AGGICTACIT CCIGGCCCTG GCAGCIGATT GGCTTCAGGC CCCCTACCTC
TATAAACTCT ACCAGCATTA CTACTICCTG GAAGGICAAA TIGCCATCCT CTATGICTGT GGCCTIGCCT CTACAGTCCT
CTITGGCCTA GIGGCCTCCT CCCTTGTGGA TIGGCTGGGT CGCAAGAATT CTTGTGTCCT CTTCTCCCTG ACTTACTCAC
TATGCTGCTT AACCAAACTC TCTCAAGACT ACTTGTGCT GCTAGTGGG CGAGCACTTG GTGGGCTGTC CACAGCCTGG
CTCTTCTCAG CCTTCGAGGN CTGGTATATC CATGAGCACG TGGAACGGGC ATGACTTTCC CTGCTGAGTG GATCCCAGCT
AACC

<u>SEQ ID NO:2162:</u> (Length of Sequence = 339 Nucleotides)

CACTGCCTT TIGTAGCTTG GGATCTAATT TGTAACACCT TGCTACCTAT GAAAAGTGG AATGTAAAAG GGAAAAAGCA
ACTTGGCATT TACTAAACTT AGGCTAACCA AAACCCTCTG TAGAGATCCT TACTAGACAT GGGTGCAACA GCAAGCATCC
CAGAGGACCC ACCACTGGGG TATGTTTTAG GCCAATGGAG CAAATTCAAA TTTGGCTAAA AGAAGAAGAA ACTCATTTAG
TATGGCAATA ATATTTGCGT TCGACACAAA GTGGCAAACC AACACATTG GCCTAAACAT GGTTCTATAT GTTATAAAACA
TACTTTACAA TTAGACTTC

SEQ ID NO:2163: (Length of Sequence = 285 Nucleotides)

CCCCGCCACC TCCAGCAGGA GCAGCTCAGT TTGTGGCTCT GGGAGCTCCG CTTTTGCAAA CCCAAAAAGG CTGTGCATTT GGAAGCCAAA CGCTCAGCAT GCGGCTGCCG AGTCTGGTTT TGTGGACAAA GCAAACTGTG GAATGGCTTC TCGGTGTCTG TATAAAGGGA CAAACGGTTG CATTCACCCT TTGTACTATA ACACCGCTTC TGCATTCGCC ATATCCGTTT TTTAACCTTT TTGTCTCCCG GGAACTTCTC ATTCGATTAT NATGTCTTCT GATGA

SEO ID NO:2164: (Length of Sequence = 296 Nucleotides)

ATGITTGTAA ATCACITCCT TITCCTACAA TATTTCTAAT AAGAAAGCTT ATAACAGCAC TITTATTGACA CCCTCGGACC CGGGGCAGGAGGG TCAGCAAGAC TCCCAGCTGG CATCAGACTG TGTCTGGCCT GCTGTCGCCA TCCCTGAGGG GTGCAGGACA GAGCCCCCATA GGGCAGAGAG GCCTCCCTGG GACCAGAGGA GGATGCTGTG CAGCCAGGCC CATCCCCAGC ACTCGAGGCC TAGGAGAGAA GGTGGGGAGA GGTGGGGGAGA GGTGGAGAGAA GCTGAGAGAA GCTGAGAGAA GCTGAGAGAA GCTGAGAGAA GCTGAGAGAA GCTGAGAGAA CAGCCC CAGCCC ACTCGAGGCC TAGGAGAGAA GCTGAGAGAA GCTGAGAGAGA GCTGAGAGAA GCTGAGAA GCTGAGAGAA GCTGAGAGAA GCTGAGAGAA GCTGAGAGAA GCTGAGAGAA GCTGAGAAA GCTGAGAGAA GCTGAGAGAA GCTGAGAGAA GCTGAGAGAA GCTGAGAGAA GCTGAGAAA GCTGAGAGAA GCTGAGAGAA GCTGAGAGAA GCTGAGAAA GCTGAGAAA GCTAGAA AGCTAGAAA GCTAGAGAA GCTAGAGAA GCTAGAGAA GCTAGAGAA GCTAGAGAA GCTAGAGAA GCTAGAGAA GCTAGAGAA GCTAGAAA AGCTAGAAA GCTAGAAA GCTAGAAA GCTAGAAA AGCTAGAAA AGCTAGAAAA GCTAGAAAA AGCTAGAAA AGCTAGAAAA GCTAGAAAA AGCTAAAA AGCTAGAAAA GCT

SEO ID NO:2165: (Length of Sequence = 310 Nucleotides)

GITTITIGIA TGITTITCAA ATAATGITIT TCTGTGTGT TTTTTTTNCT TTTTTTGGAC AGGNICICAT TCCCATTGCC CAGGGTGGAG TGCAGTGGGG CGATCTCAGC TCACTGCAGC CTTGACTTCC CAGGTTCAGA TGATTCTNCC ATCTCAGCCT CCCGAGTACC TGGGATTACA GGCACACACC ATCATGCCCG GCTAATTTTT TGTATTTTTA GCAGAGACGG GGTTTTGCCA TGTGACTCAG GCTGGTCTCG AACTCCTGGG CTCAAGAGAT CCGCCTGCCT TGGCCTCCCA AAGTGTTGGG

SEQ ID NO:2166: (Length of Sequence = 361 Nucleotides)

GATGGAAACT GGAAAAAAA TAATTIGIAA GCAACAATIT TAGATTITIT TATGGAGGAT AGAGACATIT GAATCAGATA
CCAAGAAATG TATAGTAATC ACTCACATAG AAAGATGICT AAAATGGATT TIAAATGGAT TCGGGGAAAG CAAGGTCCTG
AACAACATGC TGTACATACT ACTTATAAAT CAAAGCAAAC CACTAGCAAA CTGATGTCAG TACTAACACA GGTGGAAGTG
GGATTGTGCC GGAGGGGAGA GGTAGTNAGG GTAGACTTAT TTGTACCATT TTNATTTTTG ATATTTCTTT TATATACAGA
TACATAAGTC TGTATATACA TGTATGTCCA ATTATCTCCT T

SEO ID NO:2167: (Length of Sequence = 325 Nucleotides)

TCCTGGGCTG TGCTCTGTTT GAAGGGGGG CCCTGCTCCC CTCAGATCAG TCAGGAGGAA GATGACTAAG GGGAGGGATC
CTCTGGGTGA TGGCCTCTTC CTCCTCAGGG ACCTCTGACT GCTCTGGGCC AAAGAATCTC TTGTTTCTTC TCCGAGCCCC
AGGCAGCGGT GATTCAGCCC TGCCCAACCT GATTCTNATG ACTGCGGATG CTGTGACGGA CCCAAAGGGC AAATAGGGTC
CCAGGGTCCA GGGAGGGGC CCTGCTGAGC ACTTCCGCCC CTCACCCTGN CCAGCCCCTG CCATGAGCTC TGGGCTGGGT
CTCCG

SEO ID NO:2168: (Length of Sequence = 348 Nucleotides)

GGAGAACCGT TCGCGGAGGA AAGGCGAACT AGIGITGGGA TGGCCACCAA CTGGGGGAGC CTCTTGCAGG ATAAACAGCA
GCTAGAGGAG CTGGCACGGC AGGCCGIGGA CCGGGCCCTG GCTGAGGGAG TATTGCTGAG GACCTCACAG GAGCCCACIT
CCTCGGAGGT GGTGAGCTAT GCCCCATTCA CGCTCTTCCC CTCACTGGTC CCCAGTGCCC TGCTGGAGCA AGCCTATGCT
GTGCAGATGG ACTTCAACCT GCTAGTGGAT GCTGTCAGCC AGAACNENTG CCTTCCTGGA GCAAANTCIT TTNCAGCACC
ATCAAACAGG ATGACTTTTA CCCCTCGT

SEQ ID NO:2169: (Length of Sequence = 392 Nucleotides)

ATTITUTURA GETCCAGITT GEGIGGCAGA AACTAAGACA CIGAGCIGAT GAGAGACATE GITGCTTTTC GCCCAGGCA TITATTIATT TAITTATITA TITATTITIG TAITTITTAGI AGAGACAGAG TITCACCATG TIGGCCAGGC TEGTCTCAAA CICCIGACCI CAAATGATCC ACCCACCICG GCCTCCCAAA GIGCTGGGAT TACAAGIGTG AGCCACCATG CCCGGCCACC TGTTGCATCT TTAACAGCTG TGTTTGGAAA AGGGTGAGGA ATTGATTCAT CAATATTCAA TACTAAGCTG CAAAATCAGG AATGCAGCCA ATTGGTTTAA TTGATCAAGG CTTATAAACT CTTAAGGGAC TCTAGTGAAC TGATACAAAC TA

SEQ ID NO:2170: (Length of Sequence = 273 Nucleotides)

GITGITGITG ATGCTGITGI TGITGCTITC TGITTGITTI TCITGCAATG GICAGGTCCC ACTCTGAACT CCGGGGGGCA CCAACCTGAT GCCAGTAGGA TTGCCCTTGI ATAGGGTGIC TGACAACCCC TGITGAGGGI CTCACCCTGI TGGGTGGCAC ATGGAATAGG ACCCATTTAA TGAAGCACTI TNTCCCTTGG TGGAGGTAGT GTGCTTTNCT GGGGAAAAAC CCACTTGTCT GGGCTGCCTG GATTCCTCAG AACTACCAGG AGG

SEO ID NO:2171: (Length of Sequence = 357 Nucleotides)

GTGATGTACC CCAGCACTAG GGAATGATGT GAGTAAGACC TAATCCCTGC TCTCAGGGAG CTTATAGCCT ATGGCAGCAG
CAACACTAGT AAAAATTTAC TACTITGATA GGTGCACATC TTCCTTTGGT CAGCAATTTT CTCAAAACCA CTGTAACATT
TTACTAAAAAT GCTAAGCTTT GATTGTTTTT CAACTACTTC TTGAGAGTTT CTGCATGTAT GATAAGGGCA AGACATTACA
CTGAGGTATT GATGCTGATG AGCAGCAAGG CTCACTGGCT GGTGAAGGGA TACTGATTAG CACACCAATG TGCTGCTCTT
GAACACACAC CTCCACAAAA TTACAAATTA TCTTCCA

SEO ID NO:2172: (Length of Sequence = 381 Nucleotides)

GAAGAAGGCC CATGGAGCTA AGGCCTCAGA ACACCAAAGT CTGGACTGTC TGAGGGCACA TGCTAATAAC AGGAGGCTGG
CAAAGTGGCC AGCTCCCATG CCTTTGCATG CATTINICTT TACCTCCTGC TGCCTGGGAA CATCCTTCCA GGAGCAATCG
AGTCAACAGC ACCACAGACA CTGCTATTCC GTTGAGAAAA GTTTTATATG GAAACACATA CTGATCATGA ACACAATAAA
CAGGGAGGGA AGCTCGGGCT CAGCCAGGAA ACCTGCCACA AGGAAGATGT TTGGAACTAT CCAGGAGTAG TGTCAAACAC
TAACACCATA TTTACAAGTC TAATTTGGAA CCTGGGCCCT TTTTAAGTGC AGGAGGAAGT T

SEQ ID NO:2173: (Length of Sequence = 351 Nucleotides)

GAAGTICOGG GAGOGOCTGA AGGAGOTOGT GGICOCCAAG CACGTCATGG ATGITGTGGA CGAGGAGOTG AGCAAGCTGG GCCTGCTGGA CAACCACTCC TCGGAGTICA ATGICACCCG CAACTACCTA GACTGGCTCA CGTCCATCCC TTGGGGCAAG TACAGCAACG AGAACCTGGA CCTNGCGCGG GCACAGGCAG TGCTGGAGGA AGACCACTAC GGCATNGAGG ACGTCAAGAA ACGCATCCTG GAGTTCATTG CCGTTAGCCA GCTCCGCGGC TCCACCCAGG GCAAGATCCT CTGCTTCTAT GGGCCCCCCT GGCGTGGGTA AGACCAGCAT TGCTCTGGTC C

SEO ID NO:2174: (Length of Sequence = 308 Nucleotides)

TCATTAAATA GCTTCTATGC CACACTCTGA TTAAGCCGAC TGAGGTCCCT GGGATCTGGG TCACTGGACC GAGCTGCTCG CTCGGTGGCT CCACTGCCAG GTCCGGGCGC GCTCCCCACA GCGCTCAGTT CTGGCCCAGA CAGGGCCTGA CATCCGCCGC CTGCAGTCCC GGGGTGGCCG TCACCGTTCC ACGGCCAGNG ACTCTNCCTG CTCGTCCGGG AAGGCGATGT CGAAGATCTC CCGGTAGTNT TCCACGAAGG TAACCTCCAG GGCCCTCGGT GATGAAGGCT TCCAGGTCGT AGAAGTCC

SEQ ID NO:2175: (Length of Sequence = 403 Nucleotides)

CTTGCCCAAG GGCCTGAGCT GGTGGAGGCA GAGCAGGAGT TGGATCCAGG CCTGTNTGAG GCATCCTGCC ACCTCCATCC AGACCTGGAG CAATCCCTGA GAAGGGTGGC TACCACCAGA GATGTGGCAG CTCTGGTCTC AGGAAGCATA GCCGGAGGAT GTCCCAGGGCA ACCAAACAGC CATTCATCAG TAAGGAGCCA GAGTNAGGGC TGCTAGTTCA GCCCCGGAA GGTGGTCCAG GGGCAGCCAG TNCAGAACTC AGCAGGAGCT CAGTTCCAAC TGAGCCTGAT TTCACTCCAG TGTCCACAAG GGACATCCTG

ACCTGGAGGT CCTCGGCTAC TCACCCTGGG GCCINCTTGC ACAGCCCAGG AGCTAGCCCA GGGCTGCCTC TAAATGGTTC

SEO ID NO:2176: (Length of Sequence = 399 Nucleotides)

AGGCAACTAT TGAGGGAAGA GGCAGAAAAA GGAAAAAGGA ATGTACGTAA GGCAATTTIN CITAAAAGTA CAATAAGCTT
AATAGTGTT TAGGAAGACA AGATAAAAAT TACTCAAGGC TAGCTTGGTT CTCACTGAAT AAAAACAAAG GACTAAATAC
TGAGCTCCTT CTGTGTGGAT CTAATAATCA ATGCCTTGGT CGCTATATTG GTAATCTCTG GGGTAGTCAT CCTGGTACTC
GCCATGATAC TCATCAGGGT ATTCTGCCTG ATAATCACTA TCACTGATTT CCGAACCATT TGTTCCTGTT CCTTGGCTTC
CGTTGTGAAT GACAGGTTCT GTAGGAGCAG CACAGTATTT GGGGATCATA TACTTGCCGN CCAAGGCCAT ACAAACTCA

SEQ ID NO:2177: (Length of Sequence = 302 Nucleotides)

GGITTITATA AAAATCAGAA TITTICAAAT GCATTGGICA TITTICAGAIG CATTGGICAC ATTICATTAT TCCATATCAA AAAACTGCAT TIGITAATGI CACACAAATC TCATTGGAAA GGICTICAAG TATTGGGAAG TIGICCAGGI CACAAAGAIG AATGCIAGIT TITCAAAAATT CIACITTITA CIIGAATGCI CAAATCTIAT AATTGGIAAC CCGGICAGIT TITCITTAGI TGATAGGCIT ACTGCITTITA TGIGITGAGA ATACTIGICI GIGAAACATC CAAATCTGGA AG

SEO ID NO:2178: (Length of Sequence = 343 Nucleotides)

GGITTCACTC TCCTTGCCCA GGCTGGAGGA GCAATGTCAT GATCTTGGCT CACTGCAACC TTCTCCCTTC CAGGCTCAAT
CAATTCTCCT GCCTCAGCCT CCCGAGCAGC TGGGACTACA GGTGCGTGCC ACCATGCGCA MTAGGTTTT TTTTTGTAGA
GACAGGGTTT TGCCATGTTG CCCAGGTTGG TCTCCAACTC CTGAGCTCAA GTNATCTGCC TGANGTGCTG GGATTATAGG
TGTNAGCCAC CACATCCAGC CTCCTTTTAA TGTTTTGTTG ATTATTTATA GTGAAAGATT TAAATTCCTT TCTATTTCCT
TGTGGTATAT ATTCTATAGG CTA

SEO ID NO:2179: (Length of Sequence = 377 Nucleotides)

AGATCATCAG GAATTAGATT CTCATAAGGA ACACACAACC TAGACCCCTC AGAGGTGCAG TTCACAGTAG GGITCATGCT CCTATGAGAA CCTAATGTTG CAGCTGATCT GACAGGAGGC AGAGCTCAGC TGGIAATGCT CACTCACCTG CTGCTCACCTC CTTTCTGTGT AGCTCGGCTC CTAATAGACC TGTATGTGTC CATGGTCTGC GAGTTGGGGA CCCCTGCAGG AAGTCTTGTA AATGCATGTC AGGAAACTTA CTGTTTACAG CCACATAGTT TGTAGTAGTA AGGAAACTAG GACAATTCAA ATATTCATCA MGGGGAAAAC TGGGATAAAT TGTGGGTCAA TTTCATATGT TTCATACAGG AAAAAAG

SEO ID NO:2180: (Length of Sequence = 195 Nucleotides)

GATATTICCT TITCTCAGAA CCATAATCGA TACAAGATGC AGTGACCAAT TCATTCCTTA AAACACCTGG GCTCCTTAAG CGGCTAGAAG ACACAAGTTA CATCCAGCCC ATCAGGGAGC CAGAGGGNGA GGGGTCCCCA GCCAAGCTCT GGNCAGGCCT GCCATGGGGC AGNGCCTGAC CGTNCAGCCA GAGGT

SEO ID NO:2181: (Length of Sequence = 244 Nucleotides)

TTGGGTGGGA ACGGGCCCGG AGCGGGAGGA ACGTGACTCC CCAGAGGGAA GATGGGCATC ATACTGGGCC CAGAGCTGGG
AAGGAGTTGC TGCCAGCACA GGGTGGGCCT GGACTCCCCT CGCCCCTACC CCCAGTGGTT GTGGCTGTAG CCCTAAGCCT
GGAGAGCAGG ACCGGCCCGG GGTGTNTNGN AGGCTGCCAG GTGCCTCCCAA GGGCCCCCCAC CTGCAAGTNC
CAGC

SEO ID NO:2182: (Length of Sequence = 287 Nucleotides)

and the second

CTCCTTGAGT CTGTTGACAC GTCACATGGT CAAAGTCTCC TCATTTCAGC CAGTCTCAAC ACAAAACACC CAACAGGGAT GCACTCAACT TGTTGGTTCC ATGTGGAACT AGGTGGCAGG GCGAGAGGGA AAGTAGTAGA AGGGGGCTAT GGTGTGTCTG CATTCAGTCC CCTCACATAA AGCCACATGG ATCTAGGGGG GTATCCAAGA GCTCTGGTGG GGTCCGTGTT GCACCTAAGA CATTATAGGT CAGAGCAAGT TGCTCAGAGG GTTCCAGGCA GGGGGCT

SEO ID NO:2183: (Length of Sequence = 389 Nucleotides)

GATCCAGAGA GGGCTCCAGG TGGAGTCCCT TTTTCTGCAT AAGGGGCTGT GACCGAAGCA CAGAGGGGAA AAAAAAAAGT
GGTGGGAGCC TCCTCTGGTT TCACCTGAAG AGGGAGGTGG AAGGGCCTGA AAATTAGATT TTNTTTATAA ATAATAGATA
TTATAGGTAT ATTTNCATAT TTTTACATAA TGATGCCAAC CACAAACAAT GGACCATAAA GCACTGACCT CAGAATGATC
AATTGCAAAA TGTTTAAACC CTGGGAAGCT TTTGCTTAGG AGGGCGGATA TTCTGTGTTG ATGTTATTCT ATAGCCATAA
ACTTCCCTGA ATTTNCTGCT AATGTATCCA AGTCCAGGGA AGTCACTTAA AACTCTTCAA ATGCAGGTT

SEO ID NO:2184: (Length of Sequence = 383 Nucleotides)

GCAAGAGAAG CGGTTTGGGT CTCTGAAGGA AAGGCCAAAA CCCAGAACAA AGAAGAATCC TATGACTTCT CCAAATCCTA
TGAATATAAG TCAAACCCCT CTGCCGTTGC TGGTAATGAA ACTCCTGGGG CATCTACCAA AGGTTATCCT CCTCCTGTTG
CAGCAAAACC TACCTTTGGG CGGTCTATAC TGAAGCCCTC CACTCCCATC CCTCCTCAAG AGGGTGAGGA GGTGGGAAGA
AGCAGTGAGG AGCAAGATAA TGCTCCCAAA TCAATCCTGG GGCAAAGTCA AAATATTTGA GGAAGATGGN TCCACAAGGC
CAGGTTACAG AGGAATGCAA GGAGCTTCCA GGGAAGCACA GAATTCCAAG TTTTCGGAAA TTT

SEO ID NO:2185: (Length of Sequence = 359 Nucleotides)

CTITIAATICA CATCACAGCA GICAAGGAAG TGGGGAAAGG GGAAAAAAAT CAAGIGGCAG ATATITACAT CTAAAATICA
CATTACITGI TGGATITIGA ACATGCTACC ACAATATATA CAGTAAAATA CCICITGGGA CAATGGIACA AATITIGITI
CCITITAACIT TGCITTICIG GIACAGGIAA GATCATITIT AAATCACITI TIINCITTAA ACATGAATAC ACAAAAGAAA
TGGITAGAAG TITCCITGIT TTAAATAAGC ACAGAATGCG GGAGGITAAA AACACATTTA TAGIGCIGAA TACCAATIGG
NCATCACACT CTATACATIT TITGCTCAAA TTCCIGIAC

SEO ID NO:2186: (Length of Sequence = 337 Nucleotides)

ATAGTTATAC TCAGTGAAAT TAACAAGACC CAAAGGIGGI ATTGICTAGG AATAAAAGGG ATAATTTTIG TIGITCACAA
AAGTAACTIG TCTAGCACCA CACATCAGAA AAACACAAAA ATAGCACACT CTAGTTCTAA ACAGCTATGI CTAAAATAGA
TTATATAGIA AAACCGGTAT TATACAGCAT ATTGIGGATT TGATAAACAG ATAAATATTI GCNCTGAGTA GGCTGTTTAT
AATATAACAT TINCTTATCT ATACAGAATG AAAGCCAAAA AGTTAACTGI ATAGAGATGI GCAGAACAAC ATTAAATATT
ATGGCTCAAA AGCAGGG

SEQ ID NO:2187: (Length of Sequence = 329 Nucleotides)

GCATTINICA GCACAGATAG AGCCCTGTCC CTCCACCTAG TGCCCACTCC ATGACTGITA ATAATAACAA TAATAATAAA ACTACTGGCC AAGCACGGTG GCTCATGCCT GTAATCCCAT CACTITGGGA GGTCGAGGTG GGCAGATCAC CTGGCCCAAC GCCACCGCCT CTAGCTCCGG GCTCCCTGAG GTCCCCAGGTG CCCTINNCCGG TCCCACGGCT CCCACGNTGC CACCCTGTCC TGACTCGCCA CCTGGTCTTG TGGGCAGACT GCTGATCGAG TTCACCTCAC CCATGCCCCT GGAGGCGGG GCAGAGGGAG AACCCAGGC

<u>SEQ ID NO:2188:</u> (Length of Sequence = 335 Nucleotides)

GECCCCAGCT CCTCTTCCTG CCTCININAT GGCTTGGGCT GGAGTGGGCT CTCTGGACCT GACCGGGGGT CAGACTGTGG
GTCCCTGGGT CTCCTGCCCA CTCTNACCGG GCTTCCTCCC TCCACGCTTA GGGTCTGTCC CGGGTACTCA GTCAGGCCAG
TGGGATCTTA CCCACTTCCC TGCAAGGTGC ACCTGCCCCA GGCTCAGGCT GCCCAGCGGC TCTTCCTGGA CAGTAAGAGC
AGGGCTGGGC GCCTCTTTCC TGGCCCGGAA GCCGCAGGGG CCCCTCCTCC AGAGCCTNGG CGCAAGGAAC ACAAGGCTGC
CGCTGCTCTT CCAGG

SEO ID NO:2189: (Length of Sequence = 366 Nucleotides)

AACTGGTGGA TCAGATCGAN TTCTACTTT CTNATGAAAA CCTGGAGAAG GACGCCTTT TGCTAAAACA CGTGAGGAGG
AACAAGCTGG GATATGTGAG CNTTAAGCTA CTCACATCCT TCAAAAAGGT GAAACATCTT ACACGGGACT GGAGAACCAC
AGCACATGCT TTGAAGTATT CAGTGGTCCT TGAGTTGAAT GAGGACCACC GGAAGGTGAG GGAGGACCAC CCCCGTCCCA
CTGTTCCCCA ACGAGAACCT CCCCAGCAAG ATGCTCCTGG TCTATGATCT CTACTTGTCT CCTAAGCTGT GGGCTCTGGC
CACCCCCCAG AAGGAATGGA AGGGTGCAAG AGAAGGTGAT GGAACA

SEO ID NO:2190: (Length of Sequence = 333 Nucleotides)

CTGCGATCCA GCCTAGGCAA CAGAGTTGAG ACCCTATCTC AAAACAAACA AAACAGCCAG GCACGGTGGC TCATGCCTGT
AATCCCAGCA CTTTGGGAGG TCGAGGTGGG GGGATCACCT GAGGTCCGGA GTTCGAGACC AGACTGACCA ACATGGAGAA
AGCCCATCTC TACTAAAAAAT ACAATATTAG GGGGCGTGGT GGTGCATGCC TGTAATCCCA GCTATTTGGG AGGCTGAGGC
AGGAGAATCG CTTGAACCTG GGAGGCGGAG GTTGCAGTGA GCCATGATTG AGCCATTGCA CTACAGCCTG GGCAAGAGCA
AAACTCCGTC TTC

SEO ID NO:2191: (Length of Sequence = 284 Nucleotides)

AAGITTATAA AAGITTGATT ACIGGAAAAG TICGATCTAA TICAGAAATT TCAGGCCAAA TGAAACAGCC CCTTCAAGCA
AACATGCCTT CAATCTCTCG AGGCAGGACA ATGATTCATA TICCAGGRIGT TCGAAATAGC TCCTCAAGTA CAAGTCCTGT
TTCTAAAAAAA GGCCCACCCC TTAAGACTCC AGCCTCCAAA AGCCCTAGTG AAGGTCAAAC AGCCACCANT TCTCCTAGAG
GAGCCAAGCC ATCTGTGAAA TCAGAATTAA GCCCTGTTGC CAGG

SEO ID NO:2192: (Length of Sequence = 260 Nucleotides)

ATGACGACGE CTACCTCGAG GTCATTGGCT TCACCATGAC GTNGTTGGCC GCGCTGCAGG TGGGCGGACA CGGCGAGCGG CTGACGCAGT GTCGCGAGGT GGTGCTCACC ACATCCAAGG CCATCCCGGT GCAGGTGGAT GGCGAGCCCT GCAAGCTTTC AGCCTCACGC ATCCGCATCG CCCTGCGCAA CCAGGNCACC ATGGTGCAGA AGGCCAAGNG GCGGAGCGCC NTCCCCCTTG CACAGCGACC AGCAGCCGGT

SEO ID NO:2193: (Length of Sequence = 247 Nucleotides)

GGICICAGCA CICGCIGGGI GACCCGCGGG AGCAGGCAAA GGAGGGCICC CAAGICCGII CIGCAGCACI GGGGCAGGGA ACAGACCCAG GNICCIGGGA AICCICITCI GCCIAGCIII GCCIGCCIGC CAGAGCAGGG CCIGCGGIII GGGINCIGIN ACCNICCGGG GGCGGGGAA GGGCAAGGNA GGCGGATCIC IGAAGICCCG CCCAACIICG CINCIGATCC CCCAAGGICA GAGAGGG

SEQ ID NO:2194: (Length of Sequence = 399 Nucleotides)

CCICCATCIC CCCGGITCAA GCCATTCICG TACCICACCC TCACAAGIAG CIGGGATTAI AGGIGICCGC TACCALACCI AGCIAATTIT TGCATTGITA GCAGAGATGA GGITTCGCCA GGITGGCCAG GCIGGICITG AACTCCIGAC CTCAAGIGAT CCACCCACCI TIGHTGGCCI CCCAAAGIGC TGGAATTACA GGCAACAIGT AGCCITTGAG TCIAGCITCT TCCACIAGCC TAATTCATTT GAGATTCCAC TCGATTCTAC TTGAGATTCA TCCACATTGT TGAATGCACA TTCTTTTTTA TTTGTTCTGT AGCATTCTGT TGTGCAGCTG TGCCCCCAGTT TGTTTANCTA TTCACTCTCA GTTGTTTCCA GTTTTAATGA CAACTTCAG

SEO ID NO:2195: (Length of Sequence = 172 Nucleotides)

TCAAAGTCAG CTTCTTGACC TGCAGGGCTT CAATTTGTGG CTGACAGTTT TAACTCAGAA AATCCCTGAC TTGATTGGCT ACATAAATNA TATGINATAT AGCCATTAAG ATCATGGTTT TGGAAAGTAT TTTAATGATA CAGGAATGTG CTCTGAAATA ATAAGTGGGA CT

SEO ID NO:2196: (Length of Sequence = 398 Nucleotides)

GCAAAAAAA AAATTATTAT CTCCACTTTA CCAGTGCTGA CACTTCACCA ATGTAGGGCT CTCAGTGACT AGCCCAGGGT
CATGCACAGC CTGTTTCAGC AGCTACCTTG GACTTGAACC CAGCTCGGTC TGTCTGACTC AATGCCTATA GTCTTAACCT
TTCCAGCAGC TGCTTCTTTG TCAAACAGGT CCTCCCGCAG GTTTTCACAG CCCAGCCCCT TACTCAACAA GTATTTATTG
ACAGGCCTCA GGAACACTAG GCAAGTAGGA TAGCAATGAA CAAGATGCTG ACCTTGACCT TGACCCTGCA TCCATAGTAT
GAGCATTTTA ACTGGGGGAG GGTTTGCAAA GTTCTCTTAA ACAGTCTACT ACATGCTCTG TAAGCATTTT CTTATGGG

SEQ ID NO:2197: (Length of Sequence = 313 Nucleotides)

GTCCCTTGTG CATTGAGTGC ATCCCCGCTG GTGACTAAGC TCGCAGCAAG CGGCTACCCC CCGATCTGCA AAAGGGCCTC
TCCCTTTGTG TTCTATACAT TGTGAATCTT CCCGTCTGAA GAACGCCCAG CCTGCCCAGA CAAAGCCCCG CCTTNCCCAA
AGCAGAGGGG CTGTCTGTGT CTCCAGAAAA GGGACATCGG GGGGAGGGG GGCTCAGAAA GGAGAAGGGC TGTGATCTCC
GGTCCCTTCC CCCATCATCC TTCCTTAGAC TGATGCTTTG ACTGAATCAT CACTAGCTAT GGGCATTAAA AGG

SEQ ID NO:2198: (Length of Sequence = 360 Nucleotides)

GETCTCACTA TETTGCCCAG GCTGGTCTCA AACTCCTGTT CTCAAGCGAT CCTCCTGCCT CGGNCTACCA AGGTGCTGAG
GTTACAGGCG TGAGCACTGC ACCTGGCTAG GAAACTNAGT TTTTTCAGTG GTAGAGGCTC CTAGCCAGTG GCCAAGGGAA
AGAGAGAGTT CTGGGTTCAG GGGCTGGCAG GAAGTCAGCA AGACACCAGG GACTCGGCTC CACTGGCTGG ATCTCAGGGA
AGAGCAACTG CCACAGTGGG GACCTGGAAC ACAAAGGGAA ACTGAGGCAG CAGCTGCACC ACAGTTTACA AGTAGAAAGA
CCATGCTTGA GGACAACAGA AGTTTCACTA AGGATGCACG

SEQ ID NO:2199: (Length of Sequence = 374 Nucleotides)

TITIGGGIAG TACCCITGCC CICITCATGG CCACITCAAA GTGAAGCCAG CAAAGIGATA ATACITTATC ATITAGIATT ATCATAAAGT ATTAATACIT TGTCATAAAG TCCTCCTTGA GCCCAGGGAC CATGGAAGIC AGCTAGAAGA GCCCTGAGCA AGGAGCAAGG ACTTGGGCTT CTCCACGCTT TGCTCCTGGC TTGTTTGACC TTGACTCATT CCCCATATGT CTTTGAGGAG GCTCACAAAA TACTAAAGCT GGGAGGAAAC TTGGAGATCT ATAGGTCAAA CCTCCCCATT GGGCTGATGA GAAAATACAC GCAGGCCTAG CATGGTGCCT GCCACCATGG TGGGATCCAG TATGGTTTTA TAAA

SEO ID NO:2200: (Length of Sequence = 416 Nucleotides)

CTACTAAAAA TACAAAAATT AGCCAGGGGT GGTGGTGGGC ACCTGAAATC CCTACTCAGG AGGCTGAGGC AGAGAATCGC.
TTGAACCTGG GAGGCAGAGG TTGCAGTGAG CCGAGATCGT GCCACAGCAC TTCAGCCAGG GTGACAGAGC GAGACTCCAT
CTCAAAACAA AACAAGCAA CAACAACAA CAACAAAAAA TACCTCTTGA CTTCTAAAGA CGCAAAAGTG GCCAAAAGTG
CAATACAGTA TTGTGTTTAT TTACATCTAT TTTAAATGCA TGTGTATCTG TAAATACAAA GTGATTCGTG ACTCATTGTC

TCCTCAGTCT ATAGCATTAT TAACTITCTA GGAGCAGCAG TGGAGTAGAG TGGTACTGAA TTGGTCACAG ACTTCATCCG ATTATCAGGA TCCTGG

SEO ID NO:2201: (Length of Sequence = 315 Nucleotides)

GAAACCAATA TAAATITCAA AATAAACCAG CATACAGACC AATIGCAATI TATAGAAAAA ATAAAAATGI AGAAACATCA
CCICCICICC CCGACCCCAG TACIGAAATI ATACITCCIC AGACATACTG CCCCATCACT GGGAAGGGIG CGGACAGATT
GGGTACATIT ATAGANTATI AAATAATTAA GIAACAGAGG CACCGITTIT GCATGIATGG TCCCAAAGAC TTITCAACTT
MITITICAAC ATTACAGIIG TTAAGAATGG AAATIGAAGG AATIGTACAT ATTITCACTG GCAGTITCIT ACAGA

SEQ ID NO:2202: (Length of Sequence = 328 Nucleotides)

GCTCIGICAC TCAGCCIGGA GIGCAGIGGI GIGATCICGG CTCACIGCAA CCICIGIGIC GCAGGITCAA GCAATICICA
TGCCTCAGGC TCCIGAGIAG CIGGGATTAC AAGCATGCGC CACCATGCCC AGCIAATITT TGIATTITIA GIAGATACAG
GGTTTCGCCT TCCIGACCIC AAGCIATCCA CICGICITGG TCTCICTCAG TTCIGGGATT ACAGGIATGA GCCACCATGC
CTGGCCGGAA TATATATATT TTTTACCACT CIATTTCCAG TGCCTAGACT AAAACCCAGC ACATGGIACA CGTCATACAT
AAGGAAGG

SEO ID NO:2203: (Length of Sequence = 268 Nucleotides)

ATTITUTOTO COTOCCICAT GCCACCACTO GGACCNACCO GGT: CGG AGTGGTTTT CTGGCTTGTT TCAGCCTTTT
CAGGCTCTCT TCCATCTTCT TCACAGAGTT TAATACATCT GACACGGTTT CATAGTACTT ATGAGTGCTT TCACTGAGAG
TGCCTTCTAG CCACTGCTGA ATTATTGCTT GTTTGAGCTT ATCCTTGTGT CCGCTCTGAA GCTGGAATAA GGGCTTCANA
GCACTGTCCA CATAGGAGGA AGCTTTGG

SEO ID NO:2204: (Length of Sequence = 353 Nucleotides)

<u>SEO ID NO:2205:</u> (Length of Sequence = 265 Nucleotides)

GITTCACCAT GITGGCCAGG CIGGICTCAA ATTICINACC TCAGGIGATC CACCCCTCCT CAGCCTCCCA AAGIGITGGG ACTACAGGCG TGAGTCACTG CGCCCAGCCG TGGITTTITIT TITTITAGAAA CAGTGITTITG CCATGCTGCC CAGGCTGGTC TCAAATCCAT AGGITCAAGT GATCTCCCCA CCTCAGCCTC CCAAAGIGTC GGGACCACAG GCATGAGCCA CCATGCTTGG CCAGAAAGAA GITGTTAACA AAATG

SEO ID NO:2206: (Length of Sequence = 340 Mucleotides)

GCAAAGCITA TITTITCAGI TGIGGGCICI AGIITGGIIG GGAAACIATI TCCITAGACC TGGGICACCC CTCGGGCTCC CTTAATCICC CGCCATATGI TCTCCAGAAT CAGGGCATGG TGITCIGCCC TGGIGCGACT CAGCCCGGIIT GCTTTGCACA GACTCTGGGC CAGGGCAGGA TGICGGIGIT TGCCGGGIGIT TCGCCGGGIIG TTATCTGIGG CGCTCAGIIAT GGIGCATAGI GCTCAGACACGI GCCCTAGGIG GTGTTTAAI'I GATCTGAGAC AGACTCAGAC ACAGTGGCIC ACGTCTAITAC. TCCCAGCACT TTGGGAGGCT

SEO ID NO:2207: (Length of Sequence = 348 Nucleotides)

GIGITIGITI CICTITCCAC CATAATIGIA AGCITCCTAA GGCCICCCCA GCCCIGIGGA ATIGIGGATC AATIAAACCI CIGICCITIA TAAATAACCC AGICTGAGGC AGITCTITAT AGCAGCGIGA GAATGGACTA ATACACCICC CITCITGAGI CIGGAAGAAT AIGIGAAGGG AGATGCTAA GGACTTATIT ACAGAATGGT TCTTAAAGIG CITCGGCCAAG AACTATGTAT TINIGGAGGC TGGTAGTGIT TCAGIGAATC TGAAAACCIT TGTGACATGT GAGAAAGGTA TGCTGTCTCT GAAAGCTAAG TGTATTATGA AGGACTATA AAGGGCCA

SEQ ID NO:2208: (Length of Sequence = 154 Nucleotides)

GAATCCTGCT GTGCACATTG CTTCAGATGG CTAATTATAT CTTTGGACTG TTTGTACAAC CATTGACAAA TATACTTACT TTCATTTCTG CTAATGCAAC TGAAAAGAGC ATTCTGTAAA TTGAAGAAAA ACAAATAAAC AGNAATTAAC AACC

SEO ID NO:2209: (Length of Sequence = 352 Nucleotides)

GAGGITCAGA ATCCTCCATC CAGCATCTTC CCTGGTCACA TGGTCCCAAC CTTTTGCTCC ACCCCCTTCT CTGTTCCCCC
CGCAGTCCAT GCTCCAGCCA TCCTGACTCT GTCCCTGGAT TTCTGGCTTA CTGACACCTG AGCCTGTGCA CAGGNCCTCC
CTTCTGTATA GAGCACGCTT CCCATCTTGT GGACTTGGT GGACTCGGAG GGTTCCGAGC AGCCGTTGAG
GTGANGCTCC TATGACACCT CCNCCGTGAA GCCTNCCTCA CTTTTCCATT ACCAGTGAGG CCTGCCACAG CCTGATTTGT
ACTCTGATCC TGGCACGCAT GGAAGCCATC TT

SEO ID NO:2210: (Length of Sequence = 338 Nucleotides)

GTCTTTCCAT CAAGAGTCAA TGTATATGCA AATATAGACT TAAGAACATA AGCATCCTGG TTTAATGTTG TTGTGAGCCC
TGTGGAAATA AAATTAAACT CAGTGAATGT TTACAAATCA ATACATAGTA ATCCTATATA TGAAAGCTAA GATGTATAAG
ATGTTTATAA ATTTNCTATT AGAAAATACT GCTTTCTTAA AGGTGATTTT AAAAAGCTAG CTGATATCTG ATGGCTCAAG
CATCCAGAAA ATGTATGCAA TGATAAGNCA TTGACTAGGA TGAACAGAAA AGGGATACAG GAAAAGTCCG AACACATGAA
ATTCTAAATT AACCAAGA

SEO ID NO:2211: (Length of Sequence = 353 Nucleotides)

GTTCCTGGAG TACCCTCTTC CCCCAACCCC AGACCTGCTT TCAGAGCAAA ACTCAAGTCC CTCTTCCTCC GTGAAGCTTC
TCCCTCAGCT GAGCAGTGAT CACTTACTCA CTCTTAACCC CAATCCGCTG ACTGGGTGGG GACAGCACGT CCAGCCTTCC
CACCTCTCCT GCAGGCTTCT AGACGGAGTT TCAAAAAACTG ATGAGCCTCG ATCCAGGGCT TGAAAGAAGC CAGGGTGTAA
TCTTGTTCAT GCATGCNTCC CCAGAGNCTC GCCCAGTGCC TGGNACATAG TAGGCACTCA ATAAATGCTG AATGAGTGAA
TAGTTGAATG ATAGGTGCTC AATAAATGAA TGA

SEQ ID NO:2212: (Length of Sequence = 293 Nucleotides)

GAGAAAGGAG GCAATCTCAG TCTCGGTCTC CAAAAAGGGA TACTACTAGG GAAAGCAGAA GATCTGAATC ACTGTCCCCA AGAAGAGAAA CTTCTAGAGA GAACAAAAGA TCTCAGCCAA GAGTGAAAGA TTCTTCCCCCA GGAGAAAAAT CCAGGTCCCA GAGCAGAGAA CGAGAAAGTG ATAGAGATGG GCAGAGGAGA GAGAGAGAAA GGAGANCCAG AAAGTGGTCT AGGTCCAGAT CTCATTCTAG GTCCCCCTCA AGATGTAGAC CAAAAAGTAA GAGTTCATCA TTT

SEO ID NO:2213: (Length of Sequence = 423 Nucleotides)

NATTAACACC ACAGIGATAA ACAACTITAA GCTTATGITT CITTATAGAT CACTGGCTCA CACATAATIC AAAACCCACA CAGAAGCTAA GAGICTITAC ATTAAAAAATA TICTITCCTAA AAATCCITAC TGTATGCATC TGTCCTCAAG CAGTAAAATT TGATTATGCA CCATTTTATA ATTAATATGT CACATTTACA TAGCAAAATA ATGAAGGCAC AGCTAATACA AGCAAACTTA

AACCCTTTCT ACTTCTGAGC TGGGGGTAGG GGCACACACT TGGGATTGGT TCTTCAAGTA TATATTTTIN CCAAACATTA GCTTCAGTGA AGAGTTCTGG ATGATTTTCA CAGCTACACC CCTAAAAGCT ACATGGACAG AAAGACGTCA CAAGGCGCAA GGTACATAAC GGTGGGTACA TAT

SEO ID NO:2214: (Length of Sequence = 259 Nucleotides)

GTCATGGAGA TCCACAGCAA GTACTNGCGC TGCCTGCAGG ANCAACCTCC ACAGCGGGGC GTTCTCTGAT CGAGGCTCAG
ACTITCGAGA ACGAAGAAGC CGAGACGGTC ACCGCCATGG CCTCGCTNTC CGTGGGCGTN AAGCCCGCCG AAAAGAGACC
AGATGAGGAG CCCATGGAAG AGGAGCCGCC CCTNTAGCAC TNCCTCGAAG NTGCTGTTCT CTTGTCTGTC
TTTAAGCTCA GCCAAGAAA

SEO ID NO:2215: (Length of Sequence = 378 Nucleotides)

CACACATCCT CACCCCACAG AAACTGCTGG ACACACTGAA GAAACTGAAT AAAACAGATG AAGAAATAAG CAGTTAAAAA AATAAGTCGC CCCTCCAAAA CACGACCCA TCCCACAGCG CTCCGCAGCT TCCCACCACC GCCCGCCTCA GTTCCTTTGC GTCTGTTGCC TCCCCAGCCC TGCACGCCCT GGCTGGCACT GTTGCCGCTG CATTCTCGTG TTCAGTGATG CCCTCTTCTT GTTTGAAACA AAGAAAATA ATGCATTGTG TTTTTTTAAA AAGAGGTATC TTAATACATA GTATCCTAAA AAGAGGAGCT CATGTGGCAA TTGGTGGCAA GCAGGAGGAA ATTTCTTGGG ACTINITTAG GNIGAATT

SEO ID NO:2216: (Length of Sequence = 428 Nucleotides)

GAACCCACAC TGGGGAGAAA CCATATGAAT GTAAGGAATG TGGGAAAGCC TTCAATTATT CCAACTCATT TCAGATACAT GGAAGAACTC ACACTGAGA GAAACCCTAT GTATGTAAGG AATGTGGGAA AGCCTTCACT CAGTACTCG GCCTTAGTAT GCATGTACAGA TCTCACAGTG GAGACAAGCC CTATGAATGT AAGGAATGTG GGAAATCCTT CCTTACATCC TCACGCCTTA TTCAACATAT AAGAACTCAC ACTGGAGAGA AGCCTTTTGT ATGTGTTGAA TGTGGGAAAG CCTTTGCAGT TTCCTCAAAAT CTTAGTGGGC ATTTNAGGNA CTCACACTGN AGGAGGAAGG CCTCTGAAGT NINAGATATG TGGGGNAAGT ATTTTGGGCN ATCCCCCCAT GTCTTTAATA ATCCCCAT

SEO ID NO:2217: (Length of Sequence = 408 Nucleotides)

GICATCAGAG TICATCGIGA ACACCCIGAA IGCCGGCTCG GGGGCCTTGT CIGICACCAT IGATGGCCCC TCCAAGGIGC AGCTGGACIG ICCGGACGIAC CATGGGGCCACA CATGGGGCC AGGCCCTTCA AGGCCAAGGT CACTGGICCG AGGCTTTTCC GGAGGACACA CATGTGCGACA CATCGIGGGC AGCCCTTCA AGGCCAAGGT CACTGGICCG AGGCTTTTCC GGAGGACACA AGNITCCATT CCCCAAAGGT INITCCICAA AAINNCCAGC AAAAGGIGGG ITGACINGNG GGCCCCINGG GNITITCCCA GGGCTTTCC

SEO ID NO:2218: (Length of Sequence = 316 Nucleotides)

TTTACAGAAT ATAGCTITAT TTATAGAATC TTACAAATAA AACATTTACA GTCCACATAA GTTAATTINC TITTCTAATT
TCTTCTCATA CACCTGAGTT ATTTAAAAAA ATACTGTGAT GGAACTGCAG AACTGTAAAG GGAAATAAGA ACAATAAAAT
CCTAACCTCT CTTGCAAAAA TCAGACAACT TTGTTTTAAA GTAGATGCCC AGCATATTGC CATCTCTTTG GAAGAGGACT
TACTATACTC AGCTCTTACG NTACCCAAAC AGAGAAGCCT TCTTTTTAAA ACCCAAGGTT AAGGGCCCAG TGAAGG

SEO ID NO:2219: (Length of Sequence = 319 Micleotides)

GECTICCIGI COCACAACIT TCICACGGIG GCGCCTGGAC ACAGCAGCCA CCACAGTCCA GGCCTGCAGG GCAGGGIGTG ACCCTGCCCG GGCAGCCACC CCTCCCTGAG AAGAAGCGGG CCTCGGAGGG GGATCGITCT TTGGGCTCAG TCTCTCCCTC CICCAGIGGC TICTCCAGCC CGCACAGCGG GGAGCACCAT CAGIATCCCC TICCCAAAIN TCCITCCCGA CITTTCCAAG GCTTCAGAAG CGGCCTCACC TCINGCCAGA TAGICCAGGI GATAAACITT GIGATCGIGA AATTITGITC AAGACACIT

SEO ID NO:2220: (Length of Sequence = 343 Nucleotides)

CTGGCTAACA TGGTGAAATC CCGTCTCTAC TAAAAGTACA AAAAATTAGC TGGGCGTGGT GGTGGGCACC TGTAGCCCCA
GCTACTTGGG AGGCTGAGGC AGGAGAATGG CGTGAGGCAA CAGTGCAGCC TGGGCAACAG TGCACCTCCT CCATCTCTAC
CAGCGTCCCC TCCAGTCTGC ACGGGGCAGT CCTCCTGGGC TTGACCTCTC TGTACCCCACA GCTGGGGGCC AGGCAGCCCC
CCTCTATCCC TCCCAGCACC TACTACATCG NCCTNCACAT CCCTGATTCC TGTTGTTATG GGAAACTNTT NCCAGAGATG
GAGGTTCTCT CGGAGTATCT CGG

SEO ID NO:2221: (Length of Sequence = 373 Nucleotides)

CTCTGTCTCC CAGCCCGGAG TGCAGTAGCG CAATCTTAGC TCACTGCAGT TTTGACCTCC CAGGCTCAAA TAATCCTCCC GCCTCAGCCT CCTAAGTAGC CGAGACCACA GCTGTGCGCC ACGACATCTA GCCAATTATT TGTTTTTTGT AGAGATGAGG TCTCACTGTG TTGCTCAGGC TGGGTAGGTG TCTAACTCCT AGGCTCAAGT GATCCTCCCA CCCCAGNCTC CCAAAGTGCT GGGACTACAG GCGTGAGTCA CCGCGCCTGG CTTTGTTTAA GGCATTCTTT TTCCGCAGCA TCTGTTACCA GCAGCCTGAA GNCATTCTA TAAACAATTA TCANGGAAGA CACATGGGC AGAGACCCTA AAT

SEQ ID NO:2222: (Length of Sequence = 197 Nucleotides)

GICTCCIGIA ATTCCCCCAA ACCGGITCIT GAGGATGIGA AACCAACITA TIGGGCTCAA TCCCATTIGG TCACAGGATA CIGIACGIAT CINCCITTCC AGAGATTIGA TATCACCCAG ACACCGCCAG CATACATAAA CGIGITACCA GGITTGCCCC AGTACACAG CATATATACA CCCTTGGCCA GCCTTTC

SEO ID NO:2223: (Length of Sequence = 280 Nucleotides)

TTTTTTTTT GCATTTTAG TAGAGACGGG GTTTCACTGT GTTAGCCAGG ATGGTCTCAA TCTCCTGACC TCGTGATCCA
CCTGCCTCAG CCTCCCAAAG TGCTGGGATT ACAGGCATGA GCCACTGCGC CCGGCCAACT TTTTGCATGT TTTCTTTAAA
ATTTCTCTAC TTTTAATTGT ACTTCTAATA CAGACACTTC TGAAATCAGT TTTCACATTG CTGCAGCCTT ACCAATTTGT
AGANACTGTT TATGTGATGT TTTGATTCTT CATTTATATA

<u>SEO ID NO:2224:</u> (Length of Sequence = 388 Nucleotides)

GATTGCAGGC ATGAACCACT GCGCCCAGTC GAGTGGTAAT ATTTTGAAAG GAAACCTTTT TCTGAGCAGG TCTCAAAAGA GAGGTTAAAA TACTGAGTAG ACCATGCTGT AAACAGATGT GCTGTTATTC GGGCTTTGAT ATTCCATTTA TAAAGCACAG GCAGAGCTCA GAGTAGATTT AATGTAACTC TGAAGGGCAC TAGGATTTTN AGAATGGTAA ATAAGCATTG GCTTCAACTT AAATTCAAAT CTGCATTGGC TTGTAATAAG AGACTAGCTT GTTACTGAG CTTTNAAGCC AGTTGTTTTC TCCTATCTAG CTAGGAAAGT CCTAGATGGT ATCTACTTCC AATAAAAGGC TGTTCTGGCC AGGCGCGGTG GCTCACGC

SEO ID NO:2225: (Length of Sequence = 420 Nucleotides)

GGTCGAGGAG CCTGGGCCGG GCCGGGGGG GACTACTCCG GAGTCAGGAG GCAGCAGNGG CGGAGGACGA GGATCTCTGG CAGTCAGCGC CGCTCGGACG CCGCCGGCAC CATGGGCTGC TGCACCGGAC GCTGCTCGCT CATCTGCCTC TGCGCGCTGC AGTTGGTCTC AGCATTAGAG AGGCAGATCT TTGACTTCCT TGGTTTCCAG "IGGGCGCCTA TTCTTGGAAA TTTTCTACAC ATAATAGTTG TCATATTGGG TTTGTTTGGG ACCATTCAGT ACAGACCTCG ATACATAATG GTGGACACCG ATCTAATGAC ATTCAATATC TCTGTACATC GGTCATGGTG GAGAGAACAT GGGGCCTGGT TGTNTCAAGA AGAGTGCTGC CTTCCCTCAA GCCCCATGGC ANNGATGGAC

463

SEO ID NO:2226: (Length of Sequence = 264 Mucleotides)

GTACCTGCTC CCTGCCGGCA CCTTINITGG TGGATATITA GCTGCCCTCT ACAGTGGTTA TAACATTGAA CAGATCATGT ACCTAGGCTC GGGTTTGTNC TGTGTCGGTG CCTTGGCTGG CCTCTCCACC CAGGGAACAG CACGTCTTGG CAATGCACTG GGCATGATTG GGGTTGCTGG AGGACTGGCA GCCACCCTCG GAGTCCTAAA ACCGGGCCCA GAATTACTAG CTCAGATGTC TGGAGCGATG GCTTTGGGTG GTAC

SEO ID NO:2227: (Length of Sequence = 402 Nucleotides)

AGAGGATTGG GGCACTGGGG CAGGGGGGGCT GGCACATTCC TCAGATTCTG GCATGTCATC CTGGAAGTAC TCAGCCTGGC GGTACTGCCA CAGACGCAGG TTCCCGTCCC ACGAACTGCT GACAATCTTC TCTTCAAAGG GGTGCCAACT GACGTCACGC ACACAGGCCT TGTGGTTGGT CAGCTTCTTC ACAATGTGGC CACTTAGAAG GTCGTACACA ACCACTTTGC CAGTGGAGCA GCCACTGTAG ATGAACTGCT GGCCACTGTGG GAGAACCGGC AGCGGATGAG GGTGTGCAGC ACTCCGTGGC CCCGGTAGGT CATCAAGGAG CTGTCCCCTG GGAGCTTCAG TTTCCGCCAG GCTTTTTTTNG GGCACTTTCT GCCACCGATA GT

SEO ID NO:2228: (Length of Sequence = 394 Nucleotides)

TITAAAGTGG AAACAATGIT TITAAGAGGI GATATAAAGA AATGCCCCCA CIGIAATCCC TACCATATGI TGATTCIATG
TGGIGGGAGG GAGGGGAGAA TGATTCCITT TICTAGAATC AGAGAATTIG GAAAGTATCA AGAAAGATAA TAACAGAAAG
CATGAAATAG AGITGIGCIT TGAAGATGAA TICGATGAAA TINITATGIG AAGAGGAGIT TICCAAAGTI GCAGACCCAG
GATTCCIGGC CAGAAGCATG AAAACGITTC TITCITACIG TITCITAGGAC CIAGGCAGCA TITCITCCAT GICTGCAACA
ACATAAGAAA CAACAGCCCA AACAGCAGCA GCAACATTCA TCTGCITTGG ATCCCATGGA CAGTCATGGI GICT

SEO ID NO:2229: (Length of Sequence = 342 Nucleotides)

TITITITIAG GATGATIGAG TGITICITTA AAAATAAAAA CCCCACAAAA AAGCCAGAAC ACCCTACCCA ACCCAGCCCA GIGTAACAGG TTAGCCATTA ACACAGAATA AAGAAGGTCC CAGCCACACA CGICATTACT CGGCAGAGGG TGICCAGCCT GGICGGCCGA CGICACAGGG GAGCAGGCA TGGCACCTGC GCCACGCAGA GCACCAAGGC TGAGCATGAC CACTGGAAAT AAATAAACAT GGTGCCGACA GCATCTTTGA ATTAGTAAGA CGTTAGCACA AAACAAAAAAA GCACAACGAC TG

SEO ID NO:2230: (Length of Sequence = 357 Nucleotides)

GIGGAATGCA GCCATCACAC AGTAGTITCT GAGATTGCIT CCGTCIAGGI TITATGGGAA GATATITCCI TITCTACCAT
AGGCITCAAG GCCCTCIAAT ATCCGCITGG AAATACTACA AAAACAGIGI TICAAAACIG CICTATCAAA AGGAAGGATC
CACACIGIGA GITGAATICA CACATCACAA AGAAATCICT GAGAATTCIT CIGTCIGGGI TITATAGGAAG AAATCCCGIT
TCCAACGAAG GCCTCAAAGC GGICCATATA TCCACTIGCA GATTCIACAG AAACAATGIT TCCAAACIGC TCTATCAAGA
GGAATGITGC ACTCGGIGAG TIGAATGCAC ACATCAC

SEO ID NO:2231: (Length of Sequence = 304 Nucleotides)

AAGAGACGAG GTCTCACTTT NTNGGCCAGG TTGGTCTCAA ACCCCTGGTC ACAAACAATC CTCCAGCCTC ANCCTCCCAA
AGTGCTGGCA TTACAAGCAT GAGCCACCAT GCCCAGCTTA AGGGGGATAT TTTTTATAGAG CATCTTGCCC TGGTTCTGGA
ATTCTCTGTA GATAATACAG TTAACAGATA TTCCCCTAAG TGATTAAGAA CCTTTCCATT TGACTGATTT TNCAGAAAAG
PPTACCTATG TVACCTCAGT GGGTAGCACA ATGCCTGACA CATCTTTGV. GCTCAAATGT CTCT

SEO ID NO:2232: (Length of Sequence = 354 Mucleotides)

CCTGCCACTG AGGCAGGTGC GGCCCCAGGA CCATCACCAG GAATGCNAGG CCACCCTGGA CCAGAGGTAG GAGCCCAAGG
TCCGGCCCTT GCTCTTTGAT TGTGGGCAGC CTCCTGCCCT CTCTGGGTCT CAGTTGCCCC ATCTGCAGAG CGAGGAGGCC
CGGGCTGGTT GGTCTTGAAG GCCCCTTTTCC ATGCCGACAT CATGTCACTC TAGGCCTGGG GTTCAGTTTC CTGTGGCTGG
TGATGCTGTG GTTAAGTTTG CTTGACCCCA GCAGCCCGAG GGACTGTCTG AGTCACAGCA CAGCCCCTAT TGCGTGGCTG
CTGGTGTGTG GGGTCAATTC CAGCAGATGA ATGT

SEQ ID NO:2233: (Length of Sequence = 414 Nucleotides)

CCCAAAGCCC GCACGATGCA GGCCACTNCG ATTCCACCAA GATGGACTGT GTGTGGAGCA ACTGGAAAAG TCAGGCTATT
GACCTGTTGT ATTGGCGGGA CATCAAGCAG ACGGCCATCG TGTTTGGGAG TTTCCTGCTG CTGCTCTTCT CCCTGACCCA
GTTCAGCGTG GTGAGCGTCG TGGCCTACCT GGCCCTGGCC GCACTCTCAG CCACCATCAG TTTCCGCATC TACAAGTCTG
TTTTACAAGC AGTGCAGAAA ACCGACGAAG GCCACCCTTT CAAGGCCTAC TTGGAGCTTG AGATCANCCT TTCTCAGGAG
CAGATTCAGA AGTACACGGA CTTGCCTGCA GTTCTACGTG AACAGCACAC TTAAGGAACT NAGGAGGCTC TTCCTTGTCC
AGGACCTGGT GGAT

SEO ID NO:2234: (Length of Sequence = 394 Nucleotides)

ATAATCCGAG TGCTCCATCT TCAGTGCCAT CTGGACTCCC ACCAAGTGCA ACACCCTICA NTGTGCCTTT TGGACCAGCA CCAACAGGAA TGTATCCCTC CGTGCCTCCC ACCGGACCAC CTCCAGGACC CCCAGCACCC TTTCCTCCTT CCGGACCATC ATGTCCCCCA NCTGGTGGTC CTTATCCAGC CCCAACTGTG CCGGGCCCTG GCCCCACAGG GCATATCCTA CACCAAATAT GCCCTTTINCA GAGCTACCCA GACCATATGG TGCACCCACA GATCCAGCTG CAGNIGNTCC TTTAGGTCCA TGGGGATCCA TGTTTTINTGG ACCCTTGGGC GNCAGGAATN GGAGGGCAGT ATCCTACCCN GTAATATGGC NATATNCATN TNCA

SEO ID NO:2235: (Length of Sequence = 376 Nucleotides)

CTGATATGAT GACAATAAAG GAGTATGCTG CTGCTGTTCC GCTTTGCGTC CTCGCTACAA ACGCCTGGTG GACAACATAT
TCCCTGAAGA TCCAAAAGAT GGCCTTGTGA AAACTGATAT GGAGAAATTG ACATTTTATG CAGTATCTGC TCCAGAGAAA
CTGGATCGAA TTGGTTCTTA CCTGGCAGAA AGGTTGAGCA GGGATGTTGT CAGACATCGT TCTGGGTATG TTTTGATTGC
TATGGAGGCA CTGGACCAAC TTCTCATGGC TTGCCATTCT CAAAGCATTA AGCCATTTGT AGAAAGCTTT CTTCATATGG
TGGCAAAGCT GCTGGAATCG GGGGAACCAA AGCTTCAAGT TCTTGGAACA AATTCT

SEQ ID NO:2236: (Length of Sequence = 399 Nucleotides)

TOGCAAGAAC ACTGAAAACCC AGCCAACTTC TCCTCAGCTA GGGACCAAAA CCTTTTTGTC TGTAGTCCTT CCGAGGTTGG
AGACTCTTCT GCAGCCAAGG AAAAGGTCGC GGAGACATGC GGAGACTCCG AGGTGGAGGA GGAGTCCCCA GGAAAGCGCC
TGGACGCAGG TCTCACCAAC GGCTTTGGGG GTGCGAGGAG CGAGCAGGAG CCGGGCGGCG GCCTNGGGAG GAAGGCCACA
CCCCGACGAC GCTGTGCCTC CGAGTCCAGC ATCTCCTTCA GCAACAGCCC GCTCTGCGAC TCGAGCTTTA ATGCGCCCAA
ATNTGGGCGG GGGCAAACCG GCTCTTGTGC GACGGCACAC GCTTGGAGGA CCNCAGTNAG CTGATCTTCT GCATCGAGA

SEQ_ID_NO:2237: (Length of Sequence = 234 Nucleotides)

AAANTACTAA CATTITTAAT ACAGTCTGAT CAGATCAATT CACATCACAA GGTCAACCNG GGCTTGCTCA CATGTGNCAC
AACTGAGGNA CACAATGTCC CTACCTGCCG GCTGTCCCAC CTTCCTGGTT CCCAACAGCA TTGAAACCCC CTACTTCCCT
GACCAGACTG GCATTITTTA AAATTTTGCA TAAAACTATT TCTTCCATAG NCTTCAAACA ATCAACTAGC CAAG

SEQ ID NO:2238: (Length of Sequence = 369 Nucleotides)

ATTTAAGGCT GTACTTAACT AATTTGGGCT GAGGATGAAT ATATCAGCCA CAGCACATTA AAGAATGAGC CAAGGATTTG
TCATGGTTGG TCACITTITA AAGTATTGA TTACTGCAAC TGGAGAATGA AAAGTGTATA TTGGTGAGCG CAACCTCAGT
TTCTGAGCAC TCCTGCTCTG TGGTGAGAAT CAGACAAAAA TTCATCGGGG TGAAAAAAAA AAGGCATTAC CTGATTCACA
CCCTTGTCTT GCTAGCCCTC TTCCATTCAT TTCTCACACA GCACTTTGCT CTGTTAAATC CTCTCTCTGT CTCAGACCAT
TGCTTGCCCC TTCAAAGGGT ATGGTTCAGG CTCCTTTCAA GACATTTGG

SEO ID NO:2239: (Length of Sequence = 399 Nucleotides)

TTAATATAAT ATTCAAGTCT AGCATTTGCT ATTTACAACA AATAAATATT GCCCCTCCCC AATCAGTAAA CAAACATTTT
TTTTTTCTTT TTGCTTTTTA TACAAATATT CAATCACCC ACCCCCACCC CAAATCCTCC TTCCTCACTA ACCCCCGTCT
TGCATGGTCT CGTAAAGCCC AGGACGCAGT GGIGAATGGC ACTTGCAGTG GCATGAGATT CAACATCGAT GGGACTCAGC
TGGGACTGTC CTCACTCACC GGGIGCAGAG TCTGGTCCAT GAAGAGGGTT TCTNTCTCTG CTCCCAGGGG AGGCCTGGGG
TAAGCGGTGG GTGAGACTCC CTCACTCTCA GTTGGNCCTG ATGATGGAAT CTTTNGTGCA GCCTGAGAAA GGCTAGAGT

SEO ID NO:2240: (Length of Sequence = 388 Nucleotides)

TTTTCAGAAT TCATCTCTGA CTTTAATGGC TTAAGCAAGA ACATGGTTTC CGTGGCTCCC CCTGGACTGA ATGCTGGAGG
ATATATACTT CACAGTCTGA GGCCTGGTCC CAGGAACTGC AATCTAACAG GATGGCAAGT GGTTTTGAAA CATATAGATT
TTCAGGATGG AAGTTTGATT CTTCAGATTG TGACTCATCC GTGGAAAATA AATGGTTTAG CACCTAAATC TGTATATTCC
CATCAGTGGC TTGGCTGACT CAGTTGTAAA TAGGGTACCC TCCATCTGTC TCCCACCCAT ATGCTCCACT GTCCCCAGGG
CCTCAGTGCC TGANCCCTAG GGGGATTCGA GTTGGCTGCT GGATTCATTT CCTGCAAGCA GGCCTGCA

SEQ ID NO:2241: (Length of Sequence = 377 Nucleotides)

CTCCATTITG TCCTAGITAC TITTAAGGIA TAAGCIGAAG TCATIGATIT GAGATGITIC INCITITCIA ATATAGGIGI
TIAATGGIAC ATATITCICC CTAAGTACIG CITTAGIGGC ATCCIGCAAA TICIGACATA CIGIGGITCA TITTAATITCA
TIACAAAATA CITCITAATI TCCCTTITGA TITCCTCTIT AATTCATGGG TIACITAGAA TIGIGITATI TAATTINCAA
GIACITGGCG ATTTATCICT CTCTGTTATT CATGTCTAAT TTAATCCCAG TGTGGTCTGA GAATATATIT NGATATCAAT
AAAGCIACTC CAGCTACCTI TIGATTAATG TTATCACAGI ATATCTTTTT CTATCCT

SEO ID NO:2242: (Length of Sequence = 381 Nucleotides)

CCCACATTAA CCAAACACA ACACACATGA CAAACTCTAA GTCTCCAGAC AGACACCCC AAATAGGCAC TTGGTGTTTT
CAGCTGGGGG CTGGAGAGAT CTGGGGCTTT GGCCTCCAAA GGNAGGAGCT GCTGTCCCCA GAGAGGAGAC AACAGCTTCT
GGAGGCTCTG GGGACTCATT GGATGGGTAC TGGCTAGGTA GATGGGAAGG GGGCCTGTTT AAAGAAGACC CCCCACCCCC
ACTGCCCATT TCACCACAAC AGTGACTTGC TGGAAGTTTT GTGCCCTGCG GATTTCTGAA TATAGTGGAC AGGCATTTCT
AAAGAGCGCA TCACTGAAGG GGCAGAGGCT NGCCTTTAAA TGTGGGCTTT GCATGTTTTG G

SEO ID NO:2243: (Length of Sequence = 359 Nucleotides)

ACCATTATT AAATCAGACT GITATTCTTA ACAGTTATGT AAGTTACATG TATGITTAAG TCAGAGTATT TCACATGGAA
AAGTTTTTAA CTCCTATAGG CAAGCAAAAT CATATCACAC AATATATAAG TCGGAAGGGG ATACIGCTAA ACATTCAAAT
AAGGCAAGTA TATAAAACCA ATAAAACAAT AATGAAAAAA TTCAAGCATT CCTTTAAGAG AATTCAACAC TACAAGCTAA
ATGTACITTC TGAGTGTATT CGTATAATCA AGGCAGTGTT TCTCCTTTTA AAACATCAGG AAATGGAATA AGGCTCATTA
GTAGATACAG CTGCCCTCAA GATTTCAATT TCAGTTTGC

SEO ID NO:2244: (Length of Sequence = 362 Nucleotides)

ATATGTACTA CATTTEGTEG AATACGCATE TACAATTCTT CAAAAATAGT AAAGAGCAAA ACAAACAAAA AATAGTAGAA GCACTGGAGA AATACACTAT GGCATAAACT AGTTACGGGT GGGATGTCAC ATGGACCATA TCTACACTCT GTGGCAACCT TCTTACCTGA CTCCAAAGGA TCAGATAATC AAACAGGAAA TTATGGTAGG AAATCAGAAA ATTGAAGTAT GCATTCATAT CCTAAGCATT TTATTTTAGC TCAAAATATA AAATATTCAT CAGTTAGCCA AGCTTTGGGA TGAGAGATCA TAGCCTCCTC TTTGATAGGA GTTTCTTGTT TTCTTGATTT CATGTTTCAG AG

SEO ID NO:2245: (Length of Sequence = 333 Nucleotides)

AAGGATCIGA GCGAGTTCAG TGICATTGIG GGCAACGGGG AGATTAAGCT GCCAGTGGAG ATCAGTGGGG CCATCGAGGA
GGAGTTCACT GTGGCCCGAC TCTACATCAG CAAAATCAAA TCAGAAGTCA AGTCTGIGGI CAAGCGGIGC CGGCAGCTGG
AGAACCTCCA GGIGGAGINT CACCGCAAGA TGGAAGTNAC CGGGCGGGAG CTCTCATCCT NCCAGCTCCT CATCTCTCAG
CATGAGGCCA AGATCCGCTC GCTTACGGAA TACATGCAGA GCGIGGAGCT AAAGAAGCGG CACCTGGAAG AGTCCTATGA
CTCCTTGAGC GAT

SEQ ID NO:2246: (Length of Sequence = 347 Nucleotides)

AAACTAGCTT TGGTGGGAAC TCCCTCACCC CTGCTCCCCA CAGGAAGGCA TTAATCTATT TATGAGGGAT CTACCTGCTA
TAACCCAAAC ACCCCACCAG CCCCCATCTC CCAACACCAC CACACTGGGG ATTAAATTTC AATGTGGGAT TTGGAGAGGA
CAAATATCCA AACCATAGCA GTCTTAAAGT ATTTAAATTA GAATTTAAAT TAAAATTTAA ATTACAGTAT TTAAATTAGA
ATCATTTGTG GAGTTTCTAA AAGGTATGCA TTCCTAGGCC CCTCTCAAGT TAGATTTATG GACACTGATC CCCAGTCTGG
AATTTTAAAA CAGCAAAATC TCATACT

SEO ID NO:2247: (Length of Sequence = 357 Nucleotides)

CACAGGACAT GCTCCGTCAG CACAAGCACT CCCAAGTCAA TCTGAAAAGC AGGCAGCAGC ATTGCAGGGG ACAGGTCCTC CCCTGATCTG GGTGGTGGTC TTCTCCCACT TAAAGCACTA TATACAGGGG GAGGTCCCAG GCTGGACATC TTTACCAGGG GCTGGGAGAA AGCAGGCCGT GCTCTGTGGT CTCAGAGTCT TCCTGGCGCT CTTTGGAACC TGACAGAACA TGACCTCAGT CCCAGGCCAGC GAGTGGCAGA GAGGACTTTG TACTTGGCTG CAATAAAACA TGCCCTTCTT CGCAGAGACA CGAACAATCT CGTCTCTACC AGAGGCCTGT GAGACATCAG CTCAGGA

SEO ID NO:2248: (Length of Sequence = 327 Nucleotides)

TICTCITIAT TAATGCCIAG AAAGTCAGGT TCACCCAAGG AAGTCACTGA GGGGCCACAG CATTGAAGGG TATGGGGTTT GGAGAGATAG GAGCAGGACC CACCACTCAC GTCCAGAACC CAGGGGGCAC ACCTGGTCCA AGAGGTGGAG GCATTGGTCA CTGGAGTCAC GAGGGTCACG GAGGTCACG GGCTAGGGCT AGAGGTCGTA GCCAGTNTGC AGGGCCTGGGG AGCCCCAGGG CTGATGCCCT GGGCTGGCGT AGTACTCCAC CACCTGCCGT GGCACCCT

SEO ID NO:2249: (Length of Sequence = 404 Nucleotides)

ATTITIAATI TAGGITTGIT TITATTIAAGI TITATGITAA TICCATGCIG TGITTCAGTA AGAACAATAC AGATTCIGTA
TCTGIGGCIC CAGTCAGATA TCCAGTAGTA CAAATTAGCI TCAAGITACA CATACTGAAC AAAAGAGGIT GAGCGAGCGA
AGGAGGGGAG GAGTGAGGGG AAGGAGGTAG GGGGAGGGGG AAGGAGAAGA AACAAAAGNN TIGAACAGGC ATGCAGGCTT
TTCTTACCCA CCTTCAACGC TAACCTGCTT CAGTGGGAGA GTAAAGTAGG CAAGANIGAG CAGCCACAGA ATTGTTGAAC
TGTTACCCAG CACCATGCTT TTCAGCAACA TTTTCAGCGG AGTTTCGGAA CATTTTTTTA CCAGCAAAAA CCATTACACC
GAGT

SEQ ID NO:2250: (Length of Sequence = 275 Nucleotides)

TGCCAAATAT ATATATCIGA ACATAGIGAA AAAGIAACAT TTAAAATCAG TCAAATTATT TITAAAATIC CITIGCITAA
TAGCCATTAC TTACICACCT TITGITTITG TTTTINCCIT CAACTACIAG AGIACIGIAC TITTGCTTC ATICCITCIA
TACATICIGC CITCATCCIT AAATTGITCA ACICGATAGI GCTAATATIG GTAGATAATC TACGCTAGCT GCIGITTCIT
GTACAGAAGI TGGITGATAT CGCTGATICA CITIT

SEO ID NO:2251: (Length of Sequence = 426 Nucleotides)

GGAATAAGGA GATGAGAGCA TGCTCTGCCA ACTGGCTGGG ACCTGAATGT GCTAGGCAAG TNCCACTACA TCAGCTCAAG
AACATAAACA AAAATGTAAT TTAAAAAACA GATGGTTTAA AAAAATATCT GATAAAAATT ACCTATCCCT CTCCCTTGCT
GTGAAAATAAT TTAAATAATT TATTCTAGAT GTAAAAATAA TAATACAAAA AAGTTTGTTC AAAGACACCT GTGTCCTGTT
TGTTAAGTGT GCAGTCTGGG TCCCTTGGGG TGGAGGGAGC TGGCCAAGGA ATGGCATTGT GCAGAGGCAT ACCGGGAAGC
TCTCTGGATG CAACCCCACC TCTACCGCTT GGCAGTCAAT GACCTTGGGC ATGATGTTTC TTCACTTCTC TGAGGGCTAG
GGCTTTGATT CTGAACATGG GGGGCT

SEO ID NO:2252: (Length of Sequence = 315 Nucleotides)

GAAAAGATAA ACAAAATTAA TAGACCATTA GTGAGATTAA CCAAGACAAC AGGAAAGAAG ATCITAATAA GCTCAATTAG
CAATGAAATG NGAGCTACTA CAACTGATAC CACAGAAATA CAAAAGATCA TTCAAGGCTA CTATGAACAC CTTCACGTGC
ACAAACTAGA AAACATAGAG GAGATGGATA AATTCCTGGA ATTTTAAGAN TAATACAATG GACTTTGGGG AATCAGGAGA
AAGGGTAAGA GTGGGGTGAG GGATAAAAGA CTACACATTG CATACAGTGT ACACTTCTTG GGTGATGGGT GCGCC

SEO ID NO:2253: (Length of Sequence = 335 Nucleotides)

AGATTIATIC TCATGIACAA AGCGGICAGC CCACGGGACC ATATACGACA GITGCACAGA GICCIAGAAA AACGCATCIN
TCIAAAAGGCA ACTCAGAAAG GITAAGGCAGG TGGACCCCCT CCCCCACCCC ACAACGCACA CAGAATGAAA CGGAGAAAAA
GAGAGAAAGCC AGTGGCCGGG CTGACCCAAG AGTCCCGGCC CTATGGGGIC TCCCAAGCCC CAGGGCACAG GITGGATATGG
CCTTGAAGAG AGAGCCCTGC CAGGGCTNAG GCCAGGICTC TCACTGGCTG CAGGAATNGG TAAGGGGCTC AGGCCAAGGG
GAACACTTCA GGGGG

SEO ID NO:2254: (Length of Sequence = 380 Nucleotides)

SEO ID NO:2255: (Length of Sequence = 399 Nucleotides)

ATATAAAAAG TETTTCTGTG ATTCTNCAGA GCCCAGGAGT CAGTGCTGGT GGTTGGAGGG ACCTGCCCCC ACTGGTTCAT
TTAACCCTCT GTCTCGGTGC CCTCAGAACC TCAGCCAGAA AGGCAAGGAG GAAATCAGAG CAGGAGCCTC ATACTCTTGG
TGATCTATTC ATTCTGTGAC CTCAGGGGTC ACATATAAGG TCAGTGTTTC TCGTCCCCGC CGGATCTGCA CTGCCCAACTG
GGATTGGGTT CGAACAGCTT CATAAACATC TTCAGCATTT TGTACCATCT GCTCCCCAAT GGCCAAAAATC ACATCACCAG
GMCGCAGACC CAGCCCGGTG TGCAGGGGAGA CCCCAGGATGA CTTTATGGGA TGAGTACANC ATGCTGAACA TCGGGGAAAG

SEQ ID NO:2256: (Length of Sequence = 371 Nucleotides)

TTTTTTTTT TAACTGTAAA TGCTATTTA TTTTAAACAT TTTTGTTTAC AAAAAAAAA AAAATCAATG ATTGGTACCT
TTTTTACACT CTCAGATTCC TGAATATGGA CAGATCTTCA AAGGGAGGAA GGAGTTCTCA TATGAAATTT AAGATAGACT
GTCCTGAAGG TTGTGGGGTG GGGTTTTTTG TTGTGTTTTA ATTCGCTTTT GTTTTTAAGN CACAATAAAG CTAAAATGTC
AAGTCTCTGG GAGAGATCCC CTTAAAGTTT CAGTCAAGGA GCATATCAGA GCACAGACAA GGNGACCCCA GCCTGGTGCC
CGCCGGCCCG TCCCGGCTGC CCAGGNGTAT TTGGTAGCGC ATGGGTTGAG A

SEO ID NO:2257: (Length of Sequence = 372 Nucleotides)

AACTCTATGG CACTAATGTA TGATGGATTC ATTTCCAGAC TGTCGGCCAC GGAAGCACTT CTTCATGGCC TCTGCCCTGG
ACAGCAGCCT GTCCTCCGGG CTCCCCATGT TTTTACCAGC TTCTGCTGAG TTTCTACAAT CTTGAGCTCT GCTGAGAATT
CTTTTCCTTG AAATTCTTCT ACCTAAAAGCC CCAGCCCCCA AAAGAGCATG TCTCAGGAAC TCATTATGCC CTGAGTCAAC
AAGAACTTGT TGATAAATGG CTTAAAAGTT TTTACAAGAA GTAACTTCCC TTGGTAAGGA GTAAATAATA GCTCTGGGAA
TTTTCCAGAT AAAACTATTT CATTTCTCTG GTCAGTGGCC CCATGGGGAG AG

SEQ ID NO:2258: (Length of Sequence = 340 Nucleotides)

CTCAGCCTCC TGAGAACCTG GGATTGCAGC CTCCCGAGAA CCTGGGATTG CAGGCACCTG CTGCCATGCC CAGCGAAGAT
TTTGTATTTT TAGTGGAGAC GGGGTTTCAC CATGTTGGCC AGGCGGGTCT CAAACTCCTG ACCTCGTGAT CCACCCGCCT
TGGCCCCCCCA AAGTGCTGGG ATTACAGGGG TGAGACACCA CGCTCGGCCT TTATATATAT TTTNAGAGAG GGGTCTCAT
TTTNTTGCCC AGGCTGGTCT TGAACTCCTG GGCTCAAGCA ATCTTCCCGC CTCAGNCTCT CAAAGTGCTG GGGATTACAG
GCAATGAGCC NACCGTGNCC

SEO ID NO:2259: (Length of Sequence = 394 Nucleotides)

CCCCCCAGAT CCCACTGITA GGAGAACGCC TCTGCTAACA TTTTCTCTAT CTTGTTATCC TCTGGGAATG AGACCCACTA
AAGGGCTAGA GTGTTGCTCA GTGTGAATTC CTCTTTCTCG ACTCCATCTT CGCGGTAGCT GGGACCGCCG TTCAGTCGCC
AATATGCAGC TCTTTGTCCG CGCCCAGGAG CTACACACCT TCGAGGTGAC CGGCCAGGAA ACGGTCGCCC AGATCAAGGC
TCATGTAGCC TCACTGGAGG GCATTGCCCC GGAAGATCAA GTCGTGCTCC TGGCAGGCGC GNCCCTGGGA GGATGAGGCC
ACTCTNGGCC AGTNCGGGGT GGAGGCCCTT ACTACCCTGG AAGTAGCAAG GCCGCATGCT TINGAGGTAA AGTC

SEO ID NO:2260: (Length of Sequence = 359 Nucleotides)

TITITITITE AGATCIGAGA TICCITTAAT CAGAAGCACG TGCGTCCCAC AGTGTGCTCT TCAAGCCCCA AAGGGCACGC CTCTAGGACT GCNTCCTTAG AGCGAGGCTC GGGCTCTTGG TAAAAAAGCA TITGCTTGAT TITATITAAA CAATGGTGAA TCTTCAAGGT GCCAGTCTAC ATGCCCCAACA GTCCTCCAGG MITCAAGGNC ACAGTCACCG TCACTCAGAG ACTGCCTCAT TIMGCAAGAG AGAAAAACAG TGACCACCAC AGAGGGCAGG GAGTGACAAA GCTTGTAGGC TAATGCTGCA AAAGCCGCTA GAAACTGGGG GCCACACACA AGMGCCCANC AGGTGCGCC

SEQ ID NO:2261: (Length of Sequence = 360 Nucleotides)

TTTTTTTTT GAGACAGAGT CTCGCTCTGT CGCCAGGTTG GAATGCAGTG GTGTGATCTC AGCTCACTGC AACCTCCGCC
TCCCGGGTCC AAGCAATTCC TCTGCCTCAG CCTCCTGAGT TGCTGGGACC ACAGGCGCAC GCACCACGCC AGGCTAATTT
TTGTATTTTT AGTAGAGACG GGGTGTCACC ATATTGGCCA GGCTGGTCTC TTCGAAATCT TAAATCCAAA CATTTCTATT
CTTCTAGATC CCTTGCTCAG GCGAATCCTT TCATCTTTTC CTTATTACCTC ATCAGCATGT AAGTGTCTTG ACATCTCTCT
TCTCCTTCCC TATTACCTCT CTACTCTCTN CANTTACACG

SEO ID NO:2262: (Length of Sequence = 348 Nucleotides)

CIGICAAAAA TGIATTATAT CAATAATTIT ATCAGCAGCA TITAAGAAAT AAGAAATCAT TAGACAATAG AAGACAAACA
TGGIAATGCA GICAGGCCAG CACACAATAC ACCGITTICA TCACACACTG TAACCTGAAT CCCTGGCAAT TICCTAGAGG
TATIAACATC ATACCTTATT AAGAATTATT GGCCCCNAGG AGINGGGGGG TGGGGGGGTT GCAATCTGTC CAATCAACAT
CTGGCTCTTA CTTTCTCCCN GTAGTATTAC ATTTGTATAA TATTCTTATA GGAAACAACT CAACTCCATG TTTATAAAAG
CACCATACGG TNITTCCATC CTGTACCA

SEQ ID NO:2263: (Length of Sequence = 352 Nucleotides)

CCCCAAAAGT TGACATGGTC AATGAAGAAA TAGGCAAACA GCAAAAAGTT GCAGTCATAC ACCAAATGAA AGAAGATCAA
AGCAAAAATCC CTGAAGGAAT CCAAGTTGAC TCTGACGGGC TAATCACCAT AACAACTCCC ANTAAACTTG CCACGCTCAG
TGTTCGAGCC ATGCCCCTTC CAGAAGAAGT CACCCAGNIT CTGGAAGAAA ATAGTGANIT GATTCGTTCT ATGGAGCAGT
TGACATCCTC TTTGAATNAG GGTGAAAATA CTCACATGAT TCATCAGAAG ACCCNNENGA AAATTINGGA ATTCAAAAGGA
AAACTTINAG CAACANCTAA CAGGGNENIG AT

SEO ID NO:2264: (Length of Sequence = 381 Nucleotides)

GCITACAGIC TAGAACAAGC TITITCCAGCC CACAGCCCAG GATGGCITTG AATGIGGCCC AACACAAATT CATAAACITT
CCTAAAACAT TATGAGATCT TITIGIGAIT TGIGITTTAG TICATCAGCT ATCATTAGIG TIAGIGIATT TIGIGIGIGG
CCCAAGATAA TICITCCAAT GIGGCCCAGG GAAGCAAAAA GATTGGACAC CCCTGGICTA GAAGGAAAGG CAAATATTAA
ATAACCICAG AAAGTGATAT TACAAATTGI GGIGAGITAT AAACACACTA TCAGGIGITTA TAAAGGAAGT GAAGGAAGIG
GIGAGGAAAT TCTTATCAGG GNAGTGATAT TINANTGAAG GGCCTTAGGG GATGAGIAGG G

SEO ID NO:2265: (Length of Sequence = 301 Nucleotides)

CACTOTTOCT CONTOCTGOC TITOCACAGO AGTOAGTOTG GTOCAAGOCA COATCATOTG TOACCCAGAC TACCATAGOC ATTOTCCTAAC TEGTOCTCCC ACTTECCGTC TITATTCTGC ACACAGCAGC CTGAGTTCAT ACACACACGT GCATTCATTC ATTATTTTGCT TAAAACTGTT CAATGGCTTC CCATGGAACT TGGGAGTCTG GATATCTTCA CAAGTGTGTN GCATGGCCCA GGACCAATCT GGACACCCCT NCCTGTTTGT NCATNCATGC CTTGCACCAC TNTTTGGCCT T

SEO ID NO:2266: (Length of Sequence = 360 Nucleotides)

CGCCTGCATG CCCCACAACA ACACAACTT ATTCCTCTCC CAAACATCTG TCAGGCCTGG CCTTCCTGAG CAGGAGCTGA
GCAGGAACAG GGCCTGGCTG CCTCTCCTCT GCCACAGCTC TGACCTGGGC AAGGCTGGAA GCTGGCATCG TAATGGATGG
GGGAGTGGGT GGAGGATCTG AGGGTCCCCT GGGTAGGTTC CGATACCTTG GACAGGTGGG CCTCATCCTG ACTTAGAACT
CGGGGAGGGG CCACTCTTCC TTCCCCTTCT TCCAGCAGCA GCTCCACCAC CCTCCACCTT CTGTCCTCGA CATGTGTNCC
AGAAAACCCA GCCATGAGGG ACCGCTNTGA GGAAGGGTCT

SEO ID NO:2267: (Length of Sequence = 391 Mucleotides)

GATGGAGTCT CGCTCTGTCA CCCAGGCTGG AGTGCAGTGG CAAAAATCTCG GCTCCGGACC CCCCCAAGAC ACATATGACC CACCACCCCA TCTCTGACCA TGAGGCCACC CTGAGGTGCT GGGCCCTGGG CTTCTACCCT GCGGAGATCA CACTGACCTG GCAGCGGGAT GGGGAGGACC AGACCCAGGA CACGGAGGTC GTGGAGACCA GGCCTGCAGG GGATGGAACC TTCCAGAAGT GGGCGGCTGT GGTGGTGCCT TCTGGAGAGG AGCAGAGATA CACCTGCCAT GTGCAGCATG AGGGTCTNCC CAAGNCCCTC ACCCTGAGAA TGGGAGCTTG TCTTCCCAGC CCACCATTCC CCATCGTGGG CATNATTGCT GGNCTGGTTC T

SEO ID NO:2268: (Length of Sequence = 191 Nucleotides)

و مدينتي يري

471

TATTACACAA CIGITGITAA ATTCIAGIAA GATAAATIGA TACTAAAGAA AACAAACCCA GAAAGATCAA GIGACITGGN TCACACAACA CAGGNATTAA GANGGAAATT AGIATICTIT GITGGAATAT TITCCATTIG AATAGITACA GGAAAATITA TTIGCATATT TIACAAATTA AATGIGIATT GGACATCATA GIGGGGAAAT

SEQ ID NO:2276: (Length of Sequence = 349 Mucleotides)

TCTCCAGGTC CTGGAGGCAA CCGCAGAAAC AGAACANTGC AAATGCCAGC ATTTCCGCAG ATAAGCGTGG CCCGCCAGGT GCAAACACCC CTGACATGCA GCCGTCTGTT TAAAATCTGG TTGCCCGCTG CAGCCAGTGG AGCTCAGAGG GCTGCCTGGC GGGTAAGGAC TCCAGGCACA CAGCAACAAG TGGCTGCCAC CTCAAATCCC ACGTGGAATA TGATGGGGTC CGAGCCAGCC AGTAACTCCA NGAGGGCTGT AGTGTGTAAG TTCGGCCAGA GTTTNCAGAT ATAATANCAT TGGCCCCACG ACGTAGACCT GTGGCGGCTC AGGGTTAAGA GACGGGAGC

SEO ID NO:2277: (Length of Sequence = 182 Nucleotides)

CTTIATATAG ACTCTGGTTC TAGAAACTCG CCTGCAGCCG CTGGCTGGAC CAGCACACGC TGACGGGGCC GGACTATTTA CAGGCCCATT GCGGGCTGTA CCTTGGCCAC CTNCCGGCAC GGTGCTCAGC TGTGACGNCA AAATAAGTTA GGGCCGGCCG GGCGGGGGGG GGCGGGGACG GG

SEO ID NO:2278: (Length of Sequence = 276 Nucleotides)

GTATTATTT CCCCAAATGA AGCAAAGCAA GTACTGGGGC GGAGTCATCA GAAATACCTT GGGAGGTGGT GGGAGGGGA GTCGGGAGCA TCAGGGAAAA CCCATCTCAA CTCACGCCTC TCAGGGGTTG CGACTGGAAA MICTTGCGTT TTCCATCACT GGTGCAGAAA GAACTTCCCC AGGAATGGCC AGTGGCCTTT CGCCCGTAAC AAGGNCGCAC GCTCAGAGCA GTCTTCCTCC TGGGCTGGGT GGACGCGGAG GCGCGAAGGA AAGCCT

SEO ID NO:2279: (Length of Sequence = 193 Nucleotides)

TECACCCATE GCCCCTCCA GAGCCCCAGG GCCCCTGAGC AAGCAGGGCT CTGGCAGCAG CCAGCCCATG GAGGTGCAGG
AAGGCTATGG CTTTGGC . GGAGATGATC CCTACTCAAG TGCAGAGCCC CATGTGTCAG GTGTGAAACG GTCCCGCTCA
GGTGAGGGCG AGGTGA. 3 CCTTATGCGC AAG

SEO ID NO:2280: (Length of Sequence = 401 Nucleotides)

GTGATTITCC TGICTCCGIC TCCTGAGTAG CTGGGATTAC AGGTGCCAAC CACCACGCCC AGCTAATTIT TGIAGITITA GTGGGAGACGG TTTCGCCATG TTGGCCAGGC TGGTCTCGAA CTCCTGACCT CAGGTGATCC ATTCCCCTCG GTCTCCCAAA GTGCTGGAAT TACAGGCATG ACCCATTGCG CCCGGCCCCA CTGTTTCCTT TCTAATCGAG TGAGAAAATG GTCAGTATTT CTGTCAACAA AATTCATGAG GCCTTTTGTA CGCACAGGAC TTCAGGCCTT TCTCTCAACA ATCGCCAAAG CTGGAGGCAT CCCACAATGGA GCACAATGGA GCACAATGTT AAAAAACAGG GAATGTTTCC AGAATTNTTC TTCAAGAGTA TTTACATTTT T

SEO ID NO:2281: (Length of Sequence = 217 Nucleotides)

AGCACGGGA TIGICCAAGG GICCICCGGC GCCCAGGGCA GIGGIGGIGG CAGCACGAGT GCCCACTATG CAGTCAACAG CCAGITCACN ATGGGCGGCC CCGCCATCIC CATGGCGICG CCCATGICCA TCCCGACCAA CACCATGCAC TACGGGAGCT AGGGGCCCGN CCCGCGNAAC TNACAGCACC AGGAAACCAA ATGNATGICC CTGCCCG

SEO ID NO:2282: (Length of Sequence = 302 Nucleotides)

لولمُ للموالد الموافق الدارات إلى

CCGATGGTGA AGTGGTAAGA GGTCGATGGC CTGGGAGTTC ACTTTATTAT GAAGTAGAAA TTCTGAGNCA CGACAGCACC
TCCCAGNTTT ACACTGTAAA GTATAAAGAT GGAACAGAGC TTGANTTGAA AGAGAATGAT ATTAAGNCTT TAACTTCCTT
TAGGCAAAGG AAAGGTGGCT CAACTTCCAG TTCCCCTTCC AGACGCCGAG GGAGTCGATC AAGGTCACGC TCCCGATCCC
CCGGTCGACC ACCTAAAAGT GCCCGCCGAT CTGCTTCTGC TTTCCCACCA GGGCGACATT AA

SEQ ID NO:2283: (Length of Sequence = 314 Nucleotides)

GAAAAAGTGG AAGTCATCAC CGGGGAGGAG GCGGAGAGCA ATGTGTTACA GATGCAGTGC AAGCTGTTTG TTTTTGACAA GACCTCACAG TCCTGGGTGG AGAGAGGCCG GGGGCTGCTC AGACTCAATG ACATGGCGTC CACCGATGAC GGCACACTAC AGTCCCGACT AGTGATGCCG ACCCAGGGGA GCCTGCGACT GATCCTCAAC ACCAAGCTGT GGGCCCAGAT GCAGATCGAC AAGGCCAGCG AGAAGGAGCA TTCGCATCAC AGCCATGGAC AACGAGGACC AGGGCGTGAA GGTCTTCCTG ATCT

SEO ID NO:2284: (Length of Sequence = 262 Nucleotides)

GECGTEACAC ACGCGCCCGG CCTGTTGGAG CATTITAAAA TCTGATTCCT TTCCCCCTGA AGTTTCCGTT CAACCCTTNN CTGTGGTCAG GTTGATTNCT TTAATTGCTA AAACAAGTCA AAATTCAATA TCCATGGCAG CTGACAATTC AGACTTTGGC ATATAAAGTA AAGGGTTTAT TTTTCCATTC CTCTGTAAAT GGTGTTGINT TCACTTATTT ATAGTGCTAT GAAGCTGGTC ACCTGGGAGA ATGGCATAAC TG

SEQ ID NO:2285: (Length of Sequence = 193 Nucleotides)

GIGAGACACA GICTIGCTCI GCTGCCCAGG CTGGAGGGCA GIGTCTCGAT CTTGACTCAC TGCAGCTGAT GCCCCCTGGG
TTCAAGCGNT INTCCCACCT CAGCCTCCAA GCAGCTGGGA TTACAAACAT GNACCACCAC GGCTGGGTAA TTTTTGTGTC
TTTAGTAGAG ACGGGGNITT GCCANGTTGG CCA

SEO ID NO:2287: (Length of Sequence = 342 Nucleotides)

SEO ID NO:2288: (Length of Sequence = 343 Nucleotides)

TITTIATIGI AATGAAATIT TAAAAGGCAG TTACATTAGI TACACATATA CACAACCGAC TTAATAACIG TTAGICATAG AGAACATICA AGAAATACAA ATGATITATC CACAGCACAG TTCACATCCA TAAGAAGAAA GAGAAATGGI TAAGTACITA AACTGTCCAC TGACACCTGC TTATGAAATC TTTTCCTITC TTTCTTTTTT TAAAGGAAAC TGAGATTGIT AGATGAAGCA AGCCGTCCTG CTCCCGCACA GCCTGTGAAA CCTCCATTTT GCCACTTTCA AGGTCAGTGC CCCACAGACC CTGGGCTGTT GTTGACCATA ACACTAGCTT TGG

SEQ ID NO:2289: (Length of Sequence = 160 Nucleotides)

CGGGCCGCAA AGCTCAGCTC CTGGCGGTCC AGGCCCTGGT GGCTCTTGAT GATCAGGTCC ACGCCGGCTG CCACACCNTC CTCTAGGCCC TTCAGCGGCA NAGCGNCTCC AGCACCCTGT TGTGCTCCAT GTCCGTNAAC TGCTGCACGA AGAAGCATAT

SEO ID NO:2290: (Length of Sequence = 310 Nucleotides)

CCGACTCTAC TGAAAATACA AAATTAGCCG GGCGTGGTGA CGCATGCCTG TAATCCCAGC TACTCGGGAG GCTGAGGCAG GAGAATTGCT TGAACCCGGG AGGTGGAGGT TTGCAGTGAT CACACCACTG CACTCTAGCC TGGGTGACAA GAGCAAAACT CTGTCTCAAA AAAAAAAAA AAAACNTTAA ATGAGGTCAT GAGGGTGAGA CCCTGATCCA AGCTCATAAG TGTCCTTAGA NGTGTCCTTA GAAGTGTCCT TAGGACACTT CTTTCTAAGT NTCCTAAGTT GGGGAGCTTG CTCTCCCCCAA

SEQ ID NO:2291: (Length of Sequence = 270 Nucleotides)

CAAGACAGGG TCTCATTCTA TCTATTGCCC AGGCTGGAGT GCAGTGGTGC AATCTTGGCT CACTGCAGAC TCAACCTCCC AGGNTCAAGT GATGGAATTC CCNCAGTTTG TCTTTGACAT TAAGANGACA CCACATATAG ACGGCTGTTT GTCAGTGATT GCCCAGGNAT TCATGGATGC ATTINCTCTC ACAGAGCAGC AACTAGGGAA GGAAGCACCA ACTAATAAGC TTCTCTATGC CAAGGNTATC CCAACCTACA AAGAAGAAGT

SEO ID NO:2292: (Length of Sequence = 332 Nucleotides)

CAGITGICCT ATATTCICCA CCITCCCITG GITTCATTIC TCITCGCITC CIGAATGAGA AGIGCCIGAG ATACCITCAT
TTCICTIGAA AGIATIGATC CAAGITITAGA CAAATATCIC CCCICTIGIT GAGAGAATIC CITATATGIG AAAATACCAA
GACAITCITG ATATTTAGCA GGCACICAAA TATTIGICIC CICTITITTA GCATAATTAA GCCAGACIGA TGITIGCATT
TGAGIATCAT CAGCATGAGI AACCNITTTA ATCICTCTC CCITAACTAC TIGITCIACA CIAGAGICTA GGGICAGGGI
ACGIACAGIG AT

SEO ID NO:2293: (Length of Sequence = 255 Nucleotides)

GCACCIGACT TATGIGAGIN TCAGGCITCA ATGCCTGINT TAGAGCTACT CCITCACACA AAATAGITCA GAACATAGAG
AAGGACCAAG GITAATAAAT GATTTINATC CCAAACACTA AACATGATTG ATGGGTAGAG GCIGCCCGAA GTACTGIGTA
AAGATGGAAT CTGAGATAGA AGAATGCTGI GGICAATTAG TAATTCTTGC CCATGGAGGG ATTAGTGACA CATGCCTTGT
ATATTTGICA TCTGI

SEO ID NO:2294: (Length of Sequence = 236 Nucleotides)

GGCTTCAGAA GCTATTGGAA GATTCATATC AACTTACTAA TAATCAAGCA CTTTCATATT AAGACAATGT ATGATGTTTA GTAAAATTGA TITTNCCATA AAAGAAGTTT AAAATAAATT AGCTATTTCA AGAGNATCAT GGTTGTCAGC AAATAGAAAT GTTGTGCTTA ACTCAAATCA CAGTAATATT CTGTGGTAGT CAATTGATTT CTTTGAGCCN TTATTCTTTC ATCTGT

SEO ID NO:2295: (Length of Sequence = 308 Nucleotides)

TITTAATTIA ATCAGIAACT TIATTATAAC AAAACCIGIA TATTACCCAT TIAAACTCAT GIGIAACATT CAGIGATGIG
AGCIGIATTA AACCCAGGIA TIAGIGAAAA TITGCATIGI AAAACCIGGI AACAGIAGAC ATCIATGGGI GGICAGIAAT
TCAAGGACAC CITTTATTIT AAACAATTIT ATATAATTCA TATCAATATG CAAAATTACC ATAAAAGATA CANGGATTAA
TACATATTTA CATTTTTAGA AATAGITACT CIGAGGITGA CAGCIGICAC TITTCTAAAT ATTTACAG

SEO ID NO:2296: (Length of Sequence = 279 Nucleotides)

ACCCCICCIG GAGGCITICC CCCICCCCAG GGCITCCCTC AGGGCIACGG TGCCCGGCCA CAGITCAGIT TIGGCIACGG GCCICCACCT CCACCGCCAG ATCAGITIGC CCCICCGGGG GINICCICCT CCACCAGCCA CICCCGGGGC AGCACCICIG GCTITCCCAC CGCCICCGGC TCAGGCIGC CCGGACATGA GCAAGCCCCC GANAGCTCAG CCAGANITCC CCTATGGICA GTITCCAGGT TACGGCCAG ACTICAGIGG CTITCGGACA

SEO ID NO:2297: (Length of Sequence = 306 Nucleotides)

CTGAGAAGAA AGAGTGTGTT GTAAAGGACA ATGACTTTGA GCCCAGAGCC CTGAAAGCTA ATGGAGAAGT TATCATTGAA
ATTCCAACAA GAGCTTGTGA AGGACAAGAA AATGCTATCA AGTCCCTGGN GCATGTACAA TTTNAAGCAA CAATTGAATA
TTCCCGAAGA GGAGACCTTC ATGTCACACT TACTTCTGCT GCTGGAACTA GCACTGTGCT CTTGGCTGAA AGAGAACGGG
ATACATCTCC TAATGGCTTT AAGAATTGGG ACTTCATNGT CTGTTCACAC ATTGGGGAGA GAACCC

SEQ ID NO:2298: (Length of Sequence = 307 Nucleotides)

AGTACACCTA GTATCTITAC AGTGACTATT AAGTATTITT GAACTCAAAG TATATATTCA TCTTAAACTC CTGGAACTAT GAACCCTCCC ATGTAATTIN CTGATGAATG AAAAGGAAAA CTTTCTTTCA AATAAGTGTC ATCTGTTGCA AAAGTATGTG ATTTAAAAAC ACATGTAAAT ATAATCTTAG CTCTAATGTT TTCCTTTGGG AGTTTGGGAA AAAGCAGTTA CATTTCTCTG TTGTCTGGTT TTTATCATTT GAAAATTGGA AGGATTCATT CTGGATTGCT GAGCTGCATC AGTAGGG

SEQ ID NO:2299: (Length of Sequence = 289

GITTITAATG CATTITITIT AAAGATTAAA GIAAAATGIC TCAATTGIAA AAAATACACA CCGGGCAAAT CCTTACCTGG
MTAATAAATA TCTACATCAC AGTACAATAA AATINCINCT CTATAAAATT TAAATATGGA TTATAGICTA TCACTATCAA
AAGAAACACT ATGCTAATAT TTCCATATTA TTAAAATAAC AGGAAAAATT ACGNGCITAT TTTAGAACCT GATGCCATAG
CCGTTGGAAA GGGCAAAGAG ATTCAAATGT CGATCATCAC TCTCCATTT

SEO ID NO:2300: (Length of Sequence = 371 Nucleotides)

CACCCATTGA AAAAGCAGCC GCCCTCCTTC CCAGGAGCTG CTGAAGAGA AGCCTGCCAG AGCCTTGCCA GCAGGGACAG CCTCTTAGAT ACCAGCAGCG TCTCAGAACC CAACGTGTCC TTTGTCTCNC ACTGTGCGGA CAGCAACAGT GGTGACATAG CTGTNATCGN GGAGGTCCGG ATGGAAAACC CAAAGGAGAG TAGCAGTTCC CTGAAGACTG GGAGGCACAG CTNAGGCCAA GACAAAACCAC ACGNAACTTA CCGACTGCTG AAACGCAGGA NTCTGATCAT AGAAGCTGTC ACCAATCTTC GCTTAATCGA GAGTTTATTC ACGGTTCAGA AGATGATCAT GGATCAGGAG AAGCAGGAAG G

<u>SEO ID NO:2301:</u> (Length of Sequence = 287 Nucleotides)

ACTIGGIGIT GGGATITGIT GIGAGGITIG CIGACACCIT GACCATITIT CACTGGCIGG AAATGAAAGG AACITCCCAC TIGCICTITIG AAGGCAATIC CATTCICICC AGGGTCCITA TITCCITCCC ATATTCICIC ACACTCCCAA ACTTCIGAAG AAGGGAGCAA ACTTIGGCCA CGAGGAAGGA GINGAGCIGC CICTGTACIT GICACTGCAC CIGCACTGGT TGAATCCACC TITCCTGGGT CACGCCGCTG TGCTGGGTGG TCACAGCCTA GGACCCC

SEQ ID NO:2302: (Length of Sequence = 358 Nucleotides)

GGAACACAGG ATCCAAACTT GTCGGGGAAC TCGGAGAGAA GATCATCGTT GGCGCGGTCC TTGGTGGGCC CAAGGATGAT
GATGGGGCGA GCATAGTGCA CTTCCATCTG CGTCACTGTC TCGTAGCTCA GAACCGAGTC TTCTCGACCC TGCGATCCAG
AGCTGGAGCC CCAGTCCTTG GCCTTTAACC TTGACCACTC TCGTCGCTCA ACCCGCCGTT TGCTGGGGAT GAACCCAATG
TCGTCGGTCT CACTGTCAGA GTGGACCCGC CGTGNCTGCC ACCACTCCTC ATCACTAGCA TCGATGACAT GCAGCACATN
CCCAAAGCGG AAGTTCAAGG GCCTGGCTCA GGAAGCCG

SEO ID NO:2303: (Length of Sequence = 403 Nucleotides)

GICAGGGGCT CCAGATCATC CICCTCCAAG SGLCCCGCAG GCGCCTCCTT GGCCTCTGC TCCTGCTTGC CGCTGGCCTC
CAAGATCGTC ATGATCGAGT TAGGGATGIN AGCTTGCTCG TGGGGGGTGA AGGACCGGAC ATGGGCCAGC, AGGGGCTCCC
GGAGCTCTGG GCACTTNTCA AAGACGGCTC CCAGCTGCTG GGGCGGCANT GCAGGATGAC CTGGAAGCTC TGGGGCTTTG
TGCGCTGGCA GCACTTGATG AAGCCCTCCC ACACCTTGGG GTACTTCCAC AACTGCTTCA TGATGAGGCG GGACAGGATG

TYCATGACGG AAGCCCCCCA GGCGGGGGTA CATGGTCANG GACCTGGATG ACGGTCCTCA TGAGCAACAT GGGCAAGGGG

SEO ID NO:2304: (Length of Sequence = 376 Nucleotides)

ATCTTGCTAT GTTGCCCAGG CTGGTCTTGA ACTCCTATTC TCAAGAGAGC CTCCTGCCTC AGCCTTGTAA AGCACTGGGA
TTATAGGCAT GAACCACCGC ACCCAGCCAA GATTGCCATT TTGTATGATG AGACTGGAAG GACCCCATTG TTTCAGGATT
TTGCTACAAT ATACAAAAAAA CAATCTGTGA GACAGTGGCT GGGCTTTTTT CCTGCCTGAT TAGTTCAGTG CACATACAAC
TTGGACCAGA GGATCTGGGT TTGAATCCCA TCTCTGATAC TTCCCAAACT GAGCTGTTTT CCTTATTTGT AAAGACTAAG
ATCGCGTATG TCAAAGAGCT CTGTAAACTC TCAACACATA CAAAGTACTA CTGCTG

SEO ID NO:2305: (Length of Sequence = 354 Nucleotides)

CTGCCCAGCC TGCTCCTGGC CCCCTGGAAG CCTCCCCACA GCTGGTAATC TGGACTTAAG GATTGCTGGG CCACCGCCTC
TCTGCCTACC ACCATTCCAT ATTTAAGTGG AGCCCTACG TAGAAAGGCC CCGGGGCTTT ATTTTAGTCT CCTTTTCAGG
GATGTCGTGG GCGGGGAGG GGGTTCTTGG TGCTACAGCC CTCTCCCCAC CCCTAAAGGG ACGCCGACGC TGTTTGCTGC
CTTCACCACA TATTAGTGCT TGACCCTGGC AGGGGACCCC ATGGAAAAGA TGGGGAAGAG CAAAATACAT GGAGACGACG
CACCCINCAG GGATGCTCGC TTGGGATTCC CACG

SEO ID NO:2306: (Length of Sequence = 345 Nucleotides)

CCAAGATCCT AAGTAATTCC AAATGCCTTA GATATCAATG AAAGCTACAC ACCATTGAGA TGGGCAAAAT TCTTTCTCTA
CAAAGGGAGT AATCAAGTAA ATACCTGTCC TCTTTCAATG GACTGTTGCC TATTGAGCAT TGTGGATGAT GTGTTTTCAG
ATTTCCAGGT GAAGTTCTGA CCCTACCTGT TTGGCCAAAG ACGTAAATTG AGAGGAAAGG CCTTGGTCTT CCTGATCAAC
CAGCATTTAA CGAACAGTGG CTTAATGCAG ATCACTCAAG AGGNAGCATA GCAATGTAAA AGGAATATAA GTAGGTGTTG
GATGCCTTTT TCCTAGACCA GGAAT

SEO ID NO:2307: (Length of Sequence = 337 Nucleotides)

AACAGAATGT AAAAATACGC AAGTCAAAAC CIGGIAGAAC TGCATGGAGA AACAAATGGA TTCAATATTA TNAGICGGGA
AATTCAACGC CCTCCTATCG AAAATGGACA GATCCAGCAG GCAGAAAATT AGTAAGGACA TTGITGAGCT CIGCAATACC
ATCAATCAAC TGGATATAAT GGACATCTAT AGACTACTTC AACAACAGCA GAAGATACAT TCTTCTCAAG CTCACATGGA
ACATTCACAA AGATAGACCA CACGCAGGCC CATAAAGCAC ACCITAACAA ATTTAAAATA ATATAAATCA TACAGTGTGC
TCTCAAAACCC NCAGTGG

SEQ ID NO:2308: (Length of Sequence = 216 Nucleotides)

GAGGAGTAAA CINITITCIG AGAAGCATGC TIAGGITGIG GGACAGGAAG TGGTAAAGGC AATGCATCGI CCACAGAGGT GGATGAAGCA GINACAAAGG AATGATAATT INANCIGCIG GIGGCATCIN CACTGCTGGA GIGTATGGCA GCAATCATCI TACICICCAT CATCCTGGIG GGGGGCAGIN GIGCAGGAAA GCCACAGGGA TICGCA

SEQ ID NO:2309: (Length of Sequence = 289 Nucleotides)

GGGGCTATGA AAATACAAAA AACATTAGCA CATTCATAGT ATGTATGTGT CTACAGGCAT TINCCCAGCC CTATGAGAGT NCTGCAATTT GAGAAGTACT AAAATGTATT GTTTGGTGAC AAGAACTGCA ATAAAAAAGAT AAATGATTTN CTGAATGTTG TAGCTTAAAA ATTGCTGAGA CAAAAGACAA CTATCTGAATT AACTTAAAA ATTGCTGAGA CAAAAGACAA CTATCTGAATT AACTTAAAA ATTGCTGAGA CAAAAGACAA CTATCTGAATTT CTGCTACACAT

SEO ID NO:2310: (Length of Sequence = 359 Nucleotides)

CTENEGGCTG CCTCTCGTTG GTCAAATCCA ACCAAAAGCT AAGAGCTGGA GAGCTTGGGT GGTGCATCCA AGGAGGCTTG
CTTCCTGGGG CACAGAAGGG AGAGTGGAGA AGGATGGAAA GTGGCTCTAG GGGAGGAAAT GGAGAACATC CAGAACTTTA
TGTCACCTCT GGTGCTTGAA GGCCTTTCTC CAGGGAGACA AAAAGTTTGT NTTGGCTAAA GCTCCCTGGT TGCTCAGGAG
CCAAGGGTCA CATAATGTGC CAATGGGGT TTTTGCCTCT GAAAGCCTCT GAGGTATAAT TACTTGCAAT GNNAACATCC
CTTTTCTCTC TCTTCCTCTG CCCACCTTCC ATGCCAAGG

SEO ID NO:2311: (Length of Sequence = 324 Nucleotides)

GITINGGGCC GGCCTGGGCA ACATAGACAC CATCTCTITA AACAAACAAA CATCATTAGI TTCTACATTC TACAAGGIGA AAGACTAATT AGAAGTGAAA AATACCACTG AAATGITGGT GTACAAATGG CAGCATAATT TGATTTACAC TAGATTTTAC ACATTTTGTGT CTATTTCAAA TAGGTACTTT TACATTTTCC TTAACTGCAT CTGACACAGA GTGAATCACA GATATATGTT GGTGTCGAAA GCAGAGGTTA CTATTATTAA NCGAAAATTT TTGTGGTTTT GCAGTCATCA TATCTAATGT GGTTACAGAT TGTG

SEQ ID NO:2312: (Length of Sequence = 362 Nucleotides)

GNAGITIATA AAGCITTATI AAACATITCA AACAGCIGIG CAACGAACAC ACCAAATAAA AGCICTAGAA TAGCAGTCCA
GACGITTCAC AAGTATGGCC TCACAGTCCC ATTCCCTAGA TGGACTGCCT CCAGINCTGI NCTCTGCCTG GCCCATCTCT
CTTTCCCCTC AGGCAAGAGA GAGATGGATG GNTCAGACTG AAAGGACAGG CATGCTGATC TCCAGCAGGC AGGGGCCAGG
AGAAAGTCTC GTTTGCCAAC ACTTGTTACT GAAGCGCAGA AAAAGCAGCA AGTGACAGTC ACAAAGTCTT CCTGGGGTAT
TCTTCATAAC GTACAGTCTA TATGCGCAGG AACGAGGAAG

SEO ID NO:2313: (Length of Sequence = 449 Mucleotides)

TGTAATTITT AAATTAAGAC TECCTTAGTG AGAAAATTTC AGCAGGTGAG TTAAGGGCAC GAGGAAAGGG CCTTTGTGCA
GAAGTAATGA CATAGGCAAA TTGTCAAAGG AGAGGTTCCC TGGTGTATTT NTAGAAGAAA GTAGACCCAT GTNTCTGAAC
CCAGCACACA GTTCACTTAT GGTGGTTTTG AAATCTGCCC TGGAATTINC ATGCATCTTT TAAATTITTG GTTTATTTTTT
NCAAGAAATA AATGAAGTCT TTATTTTINC AATGAGGGCA ATGTTTATTA AGAACAGCAC ATAAGGTAGA AAAGAAGGTT
GGTTTCTAAT CTTGGTTCAT CTCCCCCACT GATCTTGAGT TTTAAAAGCA TAGAGAGCAC GATCCTTCTG TGGGGTCTCC
ACTGTCAGAG AGCCTGTNCA GATGAGCAGT CACACTGTTA CTCCACAGC

SEQ ID NO:2314: (Length of Sequence = 316 Nucleotides)

CGAGGCAAAC ACAAAGGGCT CCTTCTGCTT CTCTGACCCC ACCTGCAGCA GGTAGTGGAT AACAGCCCCT ATGGCCTCCT
TCATGACGCT CACGAGCTGC ACCTTCTGTG GCTCCTTAAG CAGTGACTGC TCACAGCGAG TGCATTCCTG GNTCCCCAAC
TCCATGAGGG CATAGCAGGC GGTCACCACA TCCTCTTTCA CCTCCGTGCC CGINTCCTCC AGTGCCAGCC GCACTTCCAC
GNACGNCAGA TTCACCAGCA GGGCCAGGAA CTTGCTCCCG GAGCTGCCCG CCGGGATCCA GTCGGAGCCG CAGGTG

SEO ID NO:2315: (Length of Sequence = 286 Nucleotides)

ATTITIATGI GIAGACAGGC TGIGGGITCC CCICACITAA ATTGAAGCIC TGITGAACTI GAGACACITA AGANICITCC
AAGINIGAAA AGIGGAGIGA AACAAAACCA TTICTAAAAC GAAAATGIGI AACINCNITC AGITITACAC AGICNAGAAA
TAAGINIVAA ACAAGITAGI CICAAACGGI TATATCITAA GGICATTITA TICCTGITAT CATTAACING ACATATCITG
GITTAGAGAG CAGCACACAA GACATTGIGI ACINITITAAT AGCTAA

SEQ ID NO:2316: (Length of Sequence = 414 Nucleotides)

المواكدة والمغيرة والسيران

AATCATAGCT TACTGTGGCC TOGATGTCCT GAGCTCAGGC GATCCTCTCC TTATAGCCTC CAGAGTAGCT GGGACTATAG
GTGCGTGCCA CCACACCCTG CTAATTINAT GTTTTGAAGA GACCGGGTCT CACTTTGTTG CCCAGGCTGG TGTCAGACTC
CTGGGCTCAA GCTAAATCAC CCACCTTGGC TTCCCAAAGT GTCGGGATTA CAGGTGTGAG CCACTGCGCC CAGCTCTGAT
TTTTTGTATTT CTTACTTAAG GCGACATACT TAGTAGCTGT GCGTCTTGGG GCAGATACCT CCCAAAGCCC CAGTTCTGTC
ATCTATAAAT AATGTAACAA CAGGGCCCCG CTCGCAGGGT TGCTGTGTGC ACATATGTGT GTGTACGTAC CCATGTGCCT
MTACGAGAAG GGCT

SEO ID NO:2317: (Length of Sequence = 166 Nucleotides)

GCAGTGACTA TTATTAACAT TACAGTACCA AGCATCCGCA AGAGACAGTC ATTTGINATT TTINATCAAG AAATAGGGCT GTTTTATACT GTTATTGACA TCAACTTTTT CCCAGTGCAT TTTTCAAAAA TATTAATAAG TTCATTCCTT TGTGCTTTTA ACTTCC

SEO ID NO:2318: (Length of Sequence = 374 Nucleotides)

TTIAITITAC ACITACAAAA GAAATGCCC ACCCTITGC CCCATTCCCC CAAAACAGTC TCTTTTACA AACATTTAAA
AATTAAAACC AAATGAAGAT AGACAAGTTA ATTTCAGTAC AATTATTTIN CAGIGTAGCT GICATAATTA GAGITTAAAT
TTCCTACAAG TGACCAATGT CCAAGIGACT TATAGGGAAA TCCTGATTAT CGGCCAAAGG AAATTCAATA TTACAAGTTA
GCAAATTCTT AGTACAAAAA TAGTCCGIGT GTTGGAACAG CTTTTCCTTG TTACATAGGT CTTAGGTCAG TCTGCTGTNA
ATACCTTAAC GNITCCGGAT TCINNICTCA CAAATG C AAATGTCACT GCTG

SEO ID NO:2319: (Length of Sequence = 380 Nucleotides)

CATCITAGIT CATGGIAATC TCCTTGGCAG CACITATTGT CITTGTGTGA GAGCAAATGA TAGAGTCATC CATTCAAGIT
AATTAAGAGC ATCTGCATTG CAAAACTGGT CACIAAATTG CTCGCCAAAT TTGAGGCTTT TTTCCTGCCA ACACAAATTA
ATTTTTTAAG TAGCAGCATT TTCAGGAGAG ACCAAATAAA GAAAGCAACA ATAAAGTTGC CTGTCTAGTG AGATGTCCCC
AAACTATCAA CTTTAAACAT ACCTTTGCCT TTNATAGTAG TTCTTCACAC AAACTGCCTT AATCAAAATG CGTGTCTCTT
GCTCTGTCAT TTTATGTTTT GGCTCTTTAG CAACCTAATT GTATGGTTAG ACAGATTCCT

SEQ ID NO:2320: (Length of Sequence = 348 Nucleotides)

GGAGITCICT TGICCACGGA GAGCAGIGIT GCAGIGIAGG GAATGCIAAA TCITACCCCA AAGGGCAAGC AGGCICCAGG
TGGCCATGAG CIGAGITGIG ACTICIGGGA ACTAATTGGG TIGGCCCCIG CIGGAGGAGC TGACAACCIG ATCAATGAGG
AGICTGACGI TGATGICCAG CICAACAACA GACACATGAI GATCCNAGGA GAAAACATGI CCAAAATCCI AAAAGCACGA
TCCATGGICA CCAGGIGCIT TAGAGATCAC TICTITNATA GGGGGGIACT ATGAAGTIAC TICCICCAAC ATTAGIGCAA
ACACAAAGTA NGAAGGIGGI GCCACACT

SEO ID NO:2321: (Length of Sequence = 330 Nucleotides)

ATCIAGACTI TNAGITCCCI GCATCIGCCA CCGIAGITIC TAGCAGGAGI AGIGGGGGGA GIAATACAGA TICINCCCIA GAAGGGGACA CIGGIAACAT GICCCACICT TGGATTAGCA GGGGIGGGIC CAGGAAGATG ATATTINCNI CITTIGCCCA CCCCCCIGGC ATTCAGCIGG ACCCAACTAG GCCATCATGA GIGGCTTCTC CCTGICATCC CCAGGGGICA TAGGATATCT ACACCGCCTT TNIGACCCCA CCCTGCACTC CCATCCITTC CTCTCTCCCC GGITCATGCC CIGCACIACA TAGCACAGCC GGGATGCTIN

SEO ID NO:2322: (Length of Sequence = 352 Nucleotides)

TTGACAAGTA AGTGTATATA TTTAAGGTGT ACAATGTGAT GCTTTGATAC ATACAGTGTG AAATGATTAC CACAGTTAGG
TTTAATAATT AACATATCCA TCATCTCACA TAGGTATGAT TTCTTATGTG TGTGGCGAGA ATCCTGAAAA TCAACTCTGA
GCACATTTCA AGTGTACAAT ACAGTATTA TGATAGTCAC CATGCTGTTA ATCAGATTGC CTACCTTGGT TAAAGTGCAG
ACTCAGGTGA AGGTCTGGAT GGAGGATCAT ACTTTAATTG ATTTAGACTC TAAAATAAAT GTATATAGTT ATTTTTGCTA
ACCTAANGAA CCTACTCATA AATGGGCTAG TG

SEO ID NO:2323: (Length of Sequence = 316 Nucleotides)

GAGACAGAGT CTCTCTCTGT CGCCCGGGCT GGAGTGCAGT GGCACAACTC AGCTCACTGC AACCTCCGCC TCCCAGATGT CCAAGTGATC AAGGGGTTC ATTTGCTCTT GGGGGATTAG GTATCATTTG GGGAGGAAGC ATGTGTTCTG TGAGGTTGTT CGGCTATGTC CAAGTGTCGT TTACTAATAG TGGAGACGGG GTTTCACCAT GTTGGCCAGG CAGGACCTCA GGTGATCTGC CCACCTCAGC CTCCCGAAGT GCTGGGATTA CAGGCATGAG TCACCACACC CGGCTTCATT TATTTTCTTA TCCATG

SEO ID NO:2324: (Length of Sequence = 300 Nucleotides)

GGGGACAGGA GGTGACCTCG CGAGCAGACG CGCGCNCCAN ACAAGCAAGC CCGCCCCGGC CTCTCGGGAG CCGTGGGGCA
GAGGCTGCGG ANCCCAGGAG GGCCGGAGCC CTCATGANIT CANTNACCTG CTTCTCCCCC TNTAGGTCTA TCAGCCACAG
TNTCTGCAAG TTTCCAAGAG CAGCAGAAAA TGAACACATT NCAGGGGCCA GTTTCATTCA AAGATGTGGC TGTGGATTTC
ACCCAGGNGG AGTGGCAGCA ACTGGACCCT GATGAGAAGA TAGCATACGG GGATGTGATG

SEO ID NO:2325: (Length of Sequence = 303 Nucleotides)

CTGTCTCAAA TAATAATGAT AATATTINCT TATGCTTACT TTACTGTAAG ATTACAGTAT ACATTACAAC ATATGCGTTT
ATTGACTGTT TATGTTATTG ATAAGGCTTC TAGTCAACAG TAGGTTACTA GTAATTAAGT TTTTGAGGAG TCAAAAGGTTA
TGTGTGGATT TTCAACTGTG GACTTTGGTG CCTCTAACCC TGTGTTGTTC AGGGGTCAAC TGTGTATTCT TTCTGTGCAA
ACATTTTTAG ATGTTATAGC CTTTAGACAT TAGAAATGGA AATTTAGTTG AACTCGAGTG TTC

SEO ID NO:2326: (Length of Sequence = 348 Nucleotides)

GITGGTCTCG TGTGGCACAT GACACAATCT CTCCCGTCCC TGGAGGCCAG CTCCCCGTG GCCAACCTCA GGCCTCCCAT
GGCATCTCAG GGCTCCTCCA GCCAGACTGG CGCCATCCAA TTAACCTGAT GGTGGCTGAG CAGCTCAGG GCAGCCTGTG
GCTGAGGGT TGGTGGTCGG GCAAGGCCAC CGTCTTGCCG AAGTCTATCA TCCAGACCTT GGCCAGGCCG GTGTGGTCGT
GCACGAAGAG GAGGGAGCTT CCTACCAC

SEO ID NO:2327: (Length of Sequence = 392 Nucleotides)

AGCTGTTTTT TCCTAGCTGC CAAGACTGTT GAGGAAGATG AGAGAATTCC AGTACTAAAG GTATTGGCAA GAGACAGTTT CTGTGGATGT TCCTCATCTG AAATTTTGAG AATGGAGGAA ATTATTCTGG ATAAGTTGAA TTGGGATCTT CACACAGCCA CACCATTGGA TTTTCTTCAT ATTTTCCATG CCATTGCAGT GTCAACTAGG CCTCAGTTAC TTTTCAGTTT GCCCAAATTG AGCCCATCTC AACATTTGGC AGTCCTTACC AMCCAACTAC TTCACTGTAT GGCCTGCAAC CAACTTCTGC AATTCAGAGG ATCCATGCTT GCTCTGGCCA TGGTTAGTCT GGAAATTGAG GAAACTCATT CCTGATTGGC TTTCTCTTTAC AA

SEQ ID NO:2328: (Length of Sequence = 256 Nucleotides)

ACGAGCACAC TCTTCACAGT GCGGCGGAAC ATCAGAAAAT GGGAGCCTTC TTCTAATGGC TGTNCTTTTC,TGTTCGCAAAA AAAAAAAAAC AAATCCTCCA AACCACACCG GATGGTTGTA AAAAGCTGCA ACGGAACCTT TGGCACCWGA TGAGAAGAGA GECCTITTAA TGCCATAGCT AGTGATGATT CANTCAAAGC ATCAGTCTAA GGAAGGATGA TGGGGGAAGG GACCNNAGAT CACAGNCCTT CTCCTT

SEO ID NO:2329: (Length of Sequence = 383 Nucleotides)

AGTAGAGACA GCATTICATT ATGITGGCCA GGCTGGTCTC GAACTCCTCA CCTCAAGTGA TCTGCCTGCC TCGGCCTCCC
AAAGTGCGGG GATTACAGGC GTGAGCACNC ATGCCTGGCC TTTTTTTTTT TTTTTTTTAA CGAAGTTATT TTTCTAGAGC
ATTCATAGTT TGTTTTTATA CAGTTAAGGT TCTCATCCAT CTGGATTTTT TGGTAAGTGT GGGGAGAATA AAATGAGGAG
CCNCTGTTTT TTTCTCCAAA TGGCATGTAT TGTCCCAACA CAATTTATTG AATCAATAAT TCATCTCTCC CATACGAATT
TAAACTATTG AACTTTCACA TCAAAATTTT GGAACTACAA AGTAGGTTTA ACAAGGTGAG AAC

SEO ID NO:2330: (Length of Sequence = 392 Nucleotides)

CEAAACCNIC TCAACCTATT CICAAACTTI AAATGGGIAA GAAGCCCACT GGICAGCATG GCAAAGCCCC AGCTCTAATA
AAAAATGCAA AAAATTGGCT GGGAGTGGAG GCGGGCGCCT GTAATCCCAG CTACTTGGAA GGTTGAGCTG GGAGAGTTGC
TTGAGICTGG GAGGCAGAGG TTGCAGTGAG CCGAGATCAC ACCACTGCAC TCCACCTTGA GCAACAGACT GAGACTCTGT
CTCAAAAAAAAA AAAAAAAANT TATGCAAAGT GTCTTTTCCA ACAAAAGTGT AATGAAGCTA GAAGTCAATA ACAGGAAAAC
CTGGGNGAAT TTGCAAGTAA GTGAAAGTTA AACAACATTC TTAACCAGTG GCTCAAAGGA GGAAATGACT GG

SEQ ID NO:2331: (Length of Sequence = 284 Nucleotides)

AAGAAAAGTA AATTCATCTT GCTCACAGTC CTTTCTGGAA GAGTTTAGAA AGCAAAGAAT TCACCGACTC AGCAGGAAGC AGAACGAGCT GTTCCTTCTT TTGACACGCA CAAGCTAATC CCCTAGAGAG TGGGGATGTG GGAAACGGAG GGTAATTAAT TCTTTGGTCA CTGGTCACT GCTGAATAGC CTTGGTCAGT TTTGGCTCTC TCCTATTTTA GGGGGAAAAA TATTTTTGTT TCTTTTTTTT AAAAAATAAA ATGTTCGCAC AATGGGAGAA AATT

SEO ID NO:2332: (Length of Sequence = 349 Nucleotides)

ATCTIAAAAA GATTITIGI ATTINCITIT GAGACIGGGI CICAGICIGI TGCCCAGGCI GGAGIGIAGC AGCCIGATCA
TGGCICAGIG CAGCCICTAC CICCCCGGC TCAGGIGATC CICCCCCTIC AGCCICCIGA GIAGCIGGGA CIACAGAGGI
GIGGCACCAT GCCCGGCIAA TITITGIATI TITIGIGGAG ATGGGGITTI GCCATGIIGC CCAGGCIAGI CITGAACICC
TGGAIGIGAG CCACIGCGIC TGGCCTATTA TITIAAATAT AGIICICITT ACIGCCAGTA GCITICATAT AACCCIAGCG
ACTAGATITA GICACCACIG CITAATICC

SEO ID NO:2333: (Length of Sequence = 353 Nucleotides)

CCACCTCTCC GITCTCTGCT TCINAACCAC AGCCGCATCC TATTTGCAGC CCTCAAGATT AAGGATGAAA ATTTGACTIT
TTAATTTTAT TATTCTTGTT CITCCTTCCT ACTTCATTAG AATCATGTTA TTGGCCTAAA ATACTGTATG TAAAGGATGC
TCIGGGGCCC ATCTGGAAGC CTGCATTCTC TGGGGATATA ATTACGCTAA GCAATTTTTC ACCAGGGACA GCATGACTTA
GCTTCTACCT GGGCATCCTC TGGCAACACA GCCCTCAGIT CITCCAAAGG GATTGGCTGC TGTCCCTTCA GGCCTTCTTC
TTGNGTGTGT GTGTGTGTG TGTGTGTGA TTC

SEQ ID NO:2334: (Length of Sequence = 279 Nucleotides)

GOGCCTTCTA CNASCTGCTG CTGCCGCNCT CATNCTGGTG GCGATGCTGC AGCTGCTCTA CCTGTCGCTG CTGTCCGGAC
TGCACAGGCA GGAGGAGCAA GACCAATATT TTAACTTCTT TCCCCCGTCC CCACGGTCCA TGGACCAGCTT CAAGGCCCAC
TCCGNACCGC GCTGGCCTCT GGAGGCGTCC TNGACGCTAG CGGCGATTAC CGCNTCTACA GGGGCCTGCT GAAGACCACC
ATNGACCCCA ACNATGTGAT CCTGGCCACG NACGCCAGC

SEO ID NO:2335: (Length of Sequence = 386 Nucleotides)

GCCTTTTGT CATGGTAGCA AAGTGGCTGC TGTGGCTCCA GGCATCACAC CCTCAATCAA GGTAGGAAGA AGAGGCCCAG
GGAGGTGTTA GCCATGCCTG TTTCTTTTAT TGGAAAAGCT TTCCCAGAAG CCCAGGTAGA CTTCCTCTC AATTTCATTG
GCCACACCTG ATCACATAGC CATCCTAAGC TGCAAAGGAG ACTGGAACAG TGAAAATCTG GATTTACAGC CTCCACAGTT
GGAGTGGCTG GAGATACAGA GTTGGGACGA CCCCTGAAAA GTGAACCAAG GTCGTCTGCA CGGCTGCCCT GGAGGGCGTG
GTGCTTGAGG TCCCTTCTAC CTCTGGGGCT TCATGGAATG ACTTGTTGCC TCCATGGAGC ACCTCT

SEO ID NO:2336: (Length of Sequence = 258 Nucleotides)

SEQ ID NO:2337: (Length of Sequence = 338 Nucleotides)

ATCICITITC CCACITCATA AAAGCAAAAT ATGIAAGACI AGCATCIGGI TITITGICCCA ATAAAAAAAT CCCACAACTI
TCAAGATATC ACTCIAGCII TCIAAAGIAG AAAGGCAATI CAGGCAACAA AAAATATIII TIAAAAATCI ATAGCCCAAA
TCACCAAAAG GIAAGGAAAG AACIITCCIA GCAAGCICIG GAGAAGACCI AATIITGGNCA TCAAAATGGA GCIITCAGAC
ACTAATCAAG GCCAITAATI AAAAAAATII TITCAGGAAA ATAAGGCAGG TIGGATCICI TITCCCACII CATAAAAGCA
AAATATGIGG CAGACICI

SEQ ID NO:2338: (Length of Sequence = 410 Nucleotides)

GEGICITECT ATECTECCTA GECTEGICTT GAACTCTTCA ACTECAGTCT TGACCTCCCA GECTCAAGTG ATCTTCTTAC
ATAGGCCTCC CAATGTGCCA GGATTATAGG CATGACCACC ATGCCAAGCT CCAGATGGTA TTCTTAATTC AGCTCACAAT
GTGCCCTCAT CAGATTGCTA GTGGCCAGGA GTGAACAACT GAGTGACTIT AAGAATCAGG ACACCAGGAA TATGTTCCTA
GAAAGTGAAG GTATGAGTGG AAAACCTGGG TTGGATTATG AACAAGGCCC ACATGTGTGC CAGAGTGGCC AGGGCAGGGA
GCAGCAGCAG GTGCTGGTGA AAGGAAGGTG GATTACTGGG GGCAATGCCT GTCTTTGTGT TATGGGTTTC TTTTGAGGGA
AGTAGATAAG

SEO ID NO:2339: (Length of Sequence = 336 Nucleotides)

ACGEGAGGA GEGECTAAGG GEGECTOGAG GAAGAGCGAA ANAGATGGAA GECTTCCGC AGAAGGCAGA GETGGGGCGT
TINTIGAGAC ATCAGTATAA CGCTCAACTC AGCAGACGCA CACAGCAGAT CCAAGAGGAG CTGGAGGCAG ACAGGCGGNT
CCTGCAGGCC CTCCTCGAGA AGGAGGACGA GAGCCAGCCC CTCCACCTGG CCAGGCGGGA GCAGGTCATG GCCGATNTCG
CCTGGNTGAA GCAGGCCCATT NAGGNGCAGC TTCAGCTGGA GCGGGCGCGG GAGGCAGAGC TGCAGATGCT TCTTGAGGGA
GGAGGGCCCAA GGAGAT

SEQ ID NO:2340: (Length of Sequence = 290 Nucleotides)

TTTTAGTAGA GATGGGGGTT TCTCCTTGIT GGTCAGGCTG GTCTCGAACT CCCGACCTCA GGTGATCCAC CTGCCTCGGC CTCCCCAAAGT GTTGGGATTA CAGGCGTGAG CACNCGCGNC CGGCCTTCAG TTTCTTCCTA GGCCGTTCTG TCACCCAAAT AGCTGCTACC CAGAGNGGCG GGGTTGACCT AGGCTGAATA TCCACTTTGT TTTTATGGAT GGCTNCCTTC CCCCATTCGGC CTTTNCCAGA ATATCCTTTC AAGTTNCANT TTCCCAGGGG AGCTCTTGGG

SEQ ID NO:2341: (Length of Sequence = 298 Nucleotides)

TTTTGCTTAT TACCCGATTT ATTAGAGAGA TCTCTAAAAA GACGGGGTGT GGCGGGGGTA GGTGGGCGAG GAACCTGGGA
TGCAAAACCAG TGTTTGGGGC CAGGAGTGGC TGTATGGTTT CANAGGCGCC CACCACTCTG GGTTTGAGGG ACACAGCACC
CTCGTCTCGG CGCTTTGGAT TNTCACGCAC CAGACCACGG GGCGGAGGAA TGGAGTGGCA TCCCTGGGGG GAGTTAAGAC
ACACGAGGTT TGCAGTTTCA TTTTGTTTCA GAATCAGTTT GGCCATAAAA ATGGGACT

SEO ID NO:2342: (Length of Sequence = 316 Nucleotides)

CCTGAACAAG GTOGTGGTGG TGTGGAATTC TCCCAAGCTG CCATCAGAGG ACCTTCTGTG GCCTGACATT GGCGTCCCCA
TCATGGTGGT CCGTACTGAG AAGAACAGTT TNAACAACCG ATTCTTACCC TGGAATGAAA TTGAGACAGA GGCCATCCTG
TCCATTGATG ACGATGCTCA CCTCCGCCAT GACGAAATCA TGTTTGGGTT CCGGGTGTGG AGAGAAGCTC GGGACCNCAT
CGTGGGCTTC CCTGGNCGTT ACCACGCATG GGACATCCCC CATCAGTCCT GGNTCTACAA CTCCAACTAC TCCTGT

SEO ID NO:2343: (Length of Sequence = 380 Nucleotides)

GGAAGAGGAG GAAGGTTGGA CCTTCATCAG ACCACTCCCT TCCCCCATCC TCCAGGAGAG GGGGCAAGGG CAACCCACCA
TCTACCCACT TACTAACCTG GTCCTAACCC CCTTACTGIG CGCGTGTGTG TGCGTGTGCG CACGCTCTGG CTGTTTGTCT
ATATGTCTAG CTCATCTAGT TCCTCTTCTT AAGGGGATGG GGGTCAGGGG CTAGGGGAGG GGGCTGAGTT TCCCCCACTTT
AGGAAGGAGGT GGGGGCTATT TCTATGCAAA TAGAAATCAG CACATTCCTC CTACTTCCCT TTCCTCCACT CCCCCCATAT
CTTTAAAGTG TGGAAGCAGA AAAGGACCTG CATTTTTCCT ACAATTCAGG AGCTGACATA

SEO ID NO:2344: (Length of Sequence = 282 Nucleotides)

GGGAATATAT TTATGCAAAT TTTATTGAAA TTTATTGTAA ATAAAGNITT TCNCAGIGGN CTAGAAAANC AGCTIGAATG
NCATTCAGCA TTTATTGAAG AAGGATGACA TCCCINCCAC TTATTGCACA AACTTGGIAG CTTTGAGACA AATACAGTAG
CACAGICCGI TTGAAGATIT GICCAAAAAA TTAGICCATA TTTTAGIGGC TCAGIGICAA GNGITCCCIC CCIGIGCCCC
CACTGTTGCT TCTGCAGIGA TACGAAGGAT GAATGCTTAA TT

SEO ID NO:2345: (Length of Sequence = 256 Nucleotides)

CTITATAGGA AGCIGCAAAA GAAATGAGCA GAGCGAGATA TITGIGGIAA GGGATACAAA GAACATACAA TIGIGIACIT GAGAGGITTC ATGGAACATT ATGACCCATC CAATGAGAC ATCAACATTA ACAACAAAAA TIANITGAGG AAGAGCAGTA TGAAAAATATT CIAATGCAGI GCIGICCAAC AGAACITICT GIGGIGATGG AAATGITCCA TATCITIGIG CIAATACAGA ATCIACCAGC CACATG

SEO ID NO:2346: (Length of Sequence = 437 Nucleotides)

GIGGAGATIG ATCCITCINI TITTIGITGC CGCTGCTGCC CTCGCGCTGG GAGCCGAGCC GGAGGGAAGG CGGTGGAGAG
ATGATTGCAG AGITGGTGAG CAGCGCTCTG GGGCTCGCCT TGTATCTCAA CACCCTGAGT GCGGATTTCT GCTATGATGA
CAGCCGTGCT ATCAAGACTA ATCAGGACCT TCTCCCAGAA ACTCCATGGA CGCACATTTT CTACAATNAT TTTTGGGGCA
CTCTTCTAAC CCACAGTGGC AGCCACAAGT CCTACCGGCC ACTCTGCACT CTTTCTTTTC GCCTGAACCA TGCCATTGGA
GGGTTGAATC CCTGGGGAGCT ACCATCTTGT CAATGTCCTG TTGCAATGCA GCAGTCACTG GTCTCTTCAC AAAGCTTCTN
CAAGATCCTC CTTTGGTGAT TGGATACTGG ACATTCA

SEC ID NO:2347: (Length of Sequence = 406 Nucleotides)

COCCEGCCCC CCTTTCCGCC GGGGGGAGAC CCCCAGGTTC AAAATGAGCC TGTTTGGGACAC AACCTCAGGT TTTGGGAACCA GTGGGACCAG CATGTTTGGC AGTGCAACTA CAGACCATCA CAATCCCATG AAGGATATTG AAGTAACATC ATCTCCTGAT GATAGCATTG GTTGTCTGTC TTTTAGCCCA CCAACCTTGC CGGGGAACTT TCTTATTGCA GGATCATGGG CTAATGATGT TOGCTGCTGG GAAGTTCAAG ACAGTGGACA GACCATTCCA AAAGCCCAGC AGATGCACAC TGGGCCTGTG CTTGATGTCT GCTGGAGTTA CGATGGGAGC AAAGTGTTTA CGGCATCGTG TGATAAAACT GCCAAAATGT GGGGACCTCA GCAGTAACCA AGCGAT

SEO ID NO:2348: (Length of Sequence = 363 Nucleotides)

GGCCTTTCAA GNAGCGGCGG ANTTCGCCGA CCGCTGTAAG GAGGTACAGC AGATCCGCGA CCAGCACCCC AGCAAAATCC CGGTGATCAT CGAGCGCTAC AAGGGTGAGA AGCAGCTGCC CGTCCTGGAC AAGACCAAGT TTTTGGTCCC GGACCATGTC AACATGAGCN AGITGGTCAA GATCATCCGG CGCCGTCTGC AGCTGAACCC CACGCAGGCC TTCTTCCTGC TGGTGAACCA GCACAGCATG GTGAGTNINT CCACGCCCAT CGCGGACATC TACGAGCAGG AGAAAGACGA GGACGGCTTC CTCTATATGG TCTACGGCTC CCAGGAAACC TTCGGCTTTC TGAGNCAGCA GTA

SEO ID NO:2349: (Length of Sequence = 332 Nucleotides)

TCCICCTACT GATGICTITC AGTAGATICA GAAGIGATIG TGGCAAACAT AGTATCTIGA AGGAAGAGAT CGIGTTITGA TTAGCATCTC CCGAGCCTAG TTTTGTGTTT ATGTTCATGG TATTGAGGAA ATAAAGATCA ATTTGGACTT CTTGCACCTG TTAATACATC CTAGTITCCTG ACTGCAGCAA AATGACTCTC AGTGCCCCTT TCTCTTCTTA GTGATTGCCT AAGATGACAG CITCATTCCC TITTAATTAT TATCCACCIT CITCCCCATC TICANTIGIT TICTCAAGIG AGGGACTIGG CCTCTACTGG GACTCCACTG GG

SEO ID NO:2350: (Length of Sequence = 339 Nucleotides)

GAGATGGAGT CTCACCCCTT CGCCCAGGCT GGAGTGCAAT GGCACGATCT CAGCTCACTG CAACCTCTTC CTCACAGGTT CAAGCAATTC TCCTGCCTCA GCCTCCCGAG TAGCTGAGAC TACAGGCGTG TGCCACCATG ACCGGCCAAT TTTTTGTACT TTTAGTAGAG ACAGGGTTTC ACCATGTTGG CCAGGCTCGC CCCGAACTCC CGACCTCATG ATCCACCTGN CTCCGCCTCC CAAAGTGCCG GGACCACAGG CATGAGNCAC CGCACCCAGA AAAAGCAAAT CTCTTAGTAT TTTTCCTCTT GTCCAAAAGG TTCTGACCAT GTTCATGAC

<u>SEQ ID NO:2351:</u> (Length of Sequence = 354 Nucleotides)

AGAAGGACCT GAGTTGTGGC CAACAACAGG CTGCAGAAAG GCAATGCCAT CCTGAAGATT TCTCAACTAA GAGTCTGCAC CCATGACAGC CCACCGAGAC CCTCGCTCCA AGTITGTGGA GAAAGGGAAC CCGCTTGGCA GCATGTGGAA AGACCCCACG ATGAGCAGCA GACACAGCAA CGCTGCCTCC TACATCTCGA CAGCATCTGT GTAAGACTCG CTAGCATCTG GTGCACACAC TGTATGAGAC AGCAACAGCC AGAACAGACA GCTTTACGTT GATGAACACA CAGACGGTGG CGCATGTTCA GAGATGCCGA GGGGACGCCG CAGTTCCCAA AATCACCTCT GGCC

SEO ID NO:2352: (Length of Sequence = 378 Nucleotides)

GITGITTGIT TAGTGGAACA CTCAAATCAA AAACAGGCTC ACGGTCTGAA TAGTCTTCTG GTCTAAGCAA CTCAGCACCA GCGCCGCCAA GGGGAGGCCG CCCTTGTCCT GGCCCCGGA AGAGACGCAG CTCCAGCCCC GACGCAGACC CCATGGCGCA CACAGGCAGG CAGAGCTCGA GGTNCAGGCG GCTGCCTTGC GGGAAGTCGC TGGGGGAGGG TCCCTNGCTG AGGCTGCACC AAGGGCTNGG GAGAGGCCCA GGAAGGGGAG AGCGAGCTGN GAGCTTGGGA TGGGAGCCGT GAGGTGGGGA TGGTTTNGCA GAGGGGCAGA GCCAAGGNCA GAGGCAAGTT CTNGGGCCCC ACAAGCTTAT GGTTGGCA

SEO ID NO:2353: (Length of Sequence = 369 Nucleotides)

و فيالفيدي CIGCCITATA TAATGIGGAT GCTGGGCACA GAGCTGTCAT CTTTGACCGA TTCCGTGGAG TGCAGGACAT TGIGGTAGGG GAAGGGACTC ATTITCTCAT CCCGTGGGTA CAGAAACCAA TTATCTTTGA CTGCCGTTCT CGACCACGTA ATGTGCCAGT CATCATTGGT AGCAAAGATT TACAGAATGT CAACATCACA CTGCGCCATCC TCTTCCGGCC TGTCGCCAGC CAGCTTCCTC
GCATCTTCAC CAGCATCGGA GAGGACTATG ATGAGGGTGT GCTGCCGTCC ATCACAACTG AGATCCTCAA GTCAGTGGTG
GCTCGCTTTG ATNCTGGAGA ACTAATCACC CAGAGAGAGC TGGTCTTCA

SEO ID NO:2354: (Length of Sequence = 363 Nucleotides)

GGAGAGGGAT TGGCATCGGC ACCATGGAGC TCCCAGGGCT TAGAGATGGA GCAAAGITGG CCTCACCTTG GGGAACCATT
CCTGCTCCTG GATACTGGAA GACATTCTGC TGCATCTNAG GATTGATTCC AGTGCCAAAC TGTCCTCCTA TGTTTCCTGT
CATGCCTCTG CTCACCATGC TGTTGCGGTT GGCCAAGGAT GCTTCAGGAT TTNCTGCTAG TTGTGAAAAC GGGCTGGTAG
AAGCAGGTGG GGTTCCTGGG ATTTGTACCA TAGTTTNGTG GATAGGGGAA TTGCTGTGGA GCACCCTGAG GAAGACGGGG
GTTNCCCATT TNAACTGGTA GTCCAGATGA GGGAGGGAGG GTT

SEO ID NO:2355: (Length of Sequence = 403 Nucleotides)

AACCAGGAAT GGAGGCCTC CTCATGTCTG AGGTAGAGTA AGACGGTGTC AGGGGGGGGA CCCGGGGGG GAGATGAGCA
CCGGCCGCAC TGGGGCATCA TCCNGGCCCA CCCGGGACGA TGGGCCGTGG GAGGGCTCAG GGCGGTGTGG TGGCCACACT
GCGAAGAATG GATTITTAAA ACACTTCATA GCCCCGANTT TNITTCAGCT CCCTCTTCGT GGACACAACT TCAGGGCTCC
CTTGTCACTG GCTTTCGGGG GTGGTCTCCC CACTTGCAGA GTCTGGTCTC CACAGGACAC CGTCCTTCCC TTCCCTTCCA
AGGGGCAGGN CCCACGNACC CTCGCCCAAA AANTAAAGGA GCTTTGTGTT TGAAAACGCC AAGGCAAGCC GTCCAAGGGA
GCT

SEO ID NO:2356: (Length of Sequence = 456 Nucleotides)

SEQ ID NO:2357: (Length of Sequence = 412 Nucleotides)

CCACCCCATE CCCAACAAGC CATATTGTCA ATAAATAAGG AATAACTGAA ACCAGACCCT TTAGGAAGAG ACAGAAATTC
CATTACCCAG GAAACCACTC AGTGAAGATG CTGATAGTTC TGATATGTTC TTATGCCCTG CCCCCTTCCC CCAAAAAAACC
ACCIGCAGAA CCAAATGTTT CTCCTCAAAG CCCATCAGCA CAGATTGATA ATAATATCAC TATCAAGCCA GGGCTAGTGC
TTCTCTACAT ACTGTACTGT CACAGGTACA AAGCAAGCCC TGGACAGATA CTGTCTCCCT GCCCCCACAA ATCCAGGGAG
GAAAAAGACC AGGGANGCTT TGATTTCCTT GGGATTTAAA CCTCATGTTC AAAAAGGNTA ATAAAGGTGC TCGTACTTGT
ATCTTCTTCC CT

SEO ID NO:2358: (Length of Sequence = 399 Nucleotides)

SEO ID NO:2359: (Length of Sequence = 352 Nucleotides)

CTICATIAAC AAGCIGCGAG AGAAGCIGGG TIGCCAGGAC GCCITCCCCG AGGIGIACGA CAAGATCIGC AAGGCCGCCA
GGACIGAGCI GGAGCCCGCC IGGAGAGACA GACACGIGIG AGIGGICAGG CATCITCCCI TCACICAAGC ITGGCIGCTI
TCCTAGATCC ACACTITCAA AGAGAAAACCC CTCCAGAACI CCCACCCIGA CAGCCCAACA CCACCITCCI CCIGGCITCC
AGGGGGGCAG CCCAGIGGAA IGGAAAGAAT GIGGGATITG GAGICAGACA AGCCTGAGIC CAGIINCCCG ITIAGAACIC
ATTAGCIGIG IGACICCIGG IGAGICCCIT AA

SEO ID NO:2360: (Length of Sequence = 359 Nucleotides)

TTTTTTTCAG CATAGTCATC TTAGCTTTAT TGAGTAAGGC ATCCCAATCT CTGCTAAGAT TCTNCTAAAT GAACGGCTGA
TTTTCCTGCC AAACTATGCA TTGGTCAAAG AGAAATCACC ACCTGGCCAC CCCATTCTGT CCCCCTACAG GACACTAAGG
GTTCTTACAG ATAAAGGGAC GATGCATTCA TGCCTGGAGA ACTAATCACA CCTGATTTCT CTGGGATCTA AANTAATGTC
AAATTTTGAT TCACTTTATG TAAAGAAAAA TCCTTTTNIT TTTNIGCAAA CCNCTTTCAA GANCAATGCT GCCCATCCCA
TGCAAGATGT TGTTGTAAGG CCANCNTCTG GTATACTAA

SEO ID NO:2361: (Length of Sequence = 437 Nucleotides)

CTCCAGGATT CCAATCCAGT CCGAACTCAA CACGAGGGG GGCACCTACA GGCTGGGGTC AATCTGGAAG ACTGCCTGTT
GTATGGCCTG GCAACTAAAA AATGTTTTT ACATTTTTAA ATGGTTAACA AAATTAAAAT AAGAGAATAT TTCATGACAT
CATCAAATTA CACGAAATGC AAATTTCAGC ATCTACAAAT ACAGTTTGAT TGGGACACAG CCACCCTCAT CCGTTTGCAG
GCTATCCCTG GCTGCTTACA GGGTCCACAT AGTCCATAAA GCCTGAGGAT ATTTACTATC TGGCCTTTTA CAGAAAAAGG
TCCCCAAAACA CTAAATCTGA AATGTTTTGC ATCAGAACCC CTTGTGGGGC TTGTTAGGAA TGCAGCTCCC TGGTCCCACA
NCCAGTCTCT GGATTCAGTA AGTCTGGAGC AGGGCCT

SEO ID NO:2362: (Length of Sequence = 317 Nucleotides)

CTTCTCTGGA TGTGCCTGGG CTTGGACTGG CTAGAATCTT TCTCTGGACT NTTGCATGTA CAGTGNCTCC ATCCTGGAGG
CAAGAGAGTT GGGACTGCT CGAATCANAG CCGTGCCCAA GATATCCCTN CTGTTGCATC GTTTGAAGCT GACGTCCTGT
GTCTNTACAC TGCTGCCACT GTTGTNTCCT CGNTCTGCTT GCTGTTGCCT CACGCCAGGN CCCGTCCTGC CGTGACANCC
TTCATCCTAC CCTTGGAACC CCAAGGCCAA GTTGGTTCAA ACTGTTGGAG AACAGAGTTG GCCTGCATCT TGGAACA

SEO ID NO:2363: (Length of Sequence = 412 Nucleotides)

GTCAGAGINI TGATAGITCI ACTGGGAGAC CACAAAATGA CATGGTCCAT CCTCCTCCTT ATCCAAAGAT GCATGGTTAA
AATAATATAG ATTAGGAATC ATCGTTACCT CCAAACAGIT AATTCAATTC AAATTTTTAG CCCAGACTGG TTTTTAAAGA
CATTTTCTGC CAAAATTTTT TGGAAGTAAA CACATTAAGG GTAGGTGTGG AGAACGATTA ATGGATTCAT TTTTATACTC
ACATCTGTTT TGGAAATATA TTTTATGCAA TAAAGCATAA ACTAACAGGT ATACTTATAA ATGTCTGGTT TTAGAAACAC
TAAAAGATCT CCAATCTTAG GAGGCCTTAA TTTGAAACTC TGCTTTTATT TGCCTGAACT AGTGGCTAAC CTGTNTAGGC
ATCTCACGAG GG

SEO ID NO:2364: (Length of Sequence = 334 Nucleotides)

GAAATGATIT AATATTAGGA AAGGCAAGIN CCTCGAGACA TITATITAAG CTAATCIGIC CITGATITIT GACITCAGA
TTCATTACAC (YAGCCACAT TAGCCIGCAC CATTAAAAAC AITGATICAA CCTCTCTTAT TGGCATACAC AATACATCIC
CCTTGITCAC TACTCTATCC TCAGCITGGT ATTTCTCTAG CACAGAAGAA TGGICCAGIA GATATGCTGA AGAAATACCT
GAATGCATAA ATAAATAAGA AAATGAGAGA CTGAATGANI CAATTAATAC CTCAAGIGIT ACCCINGATA AGGITCTAGA
GAGGGGAGGT TCTA

SEQ ID NO:2365: (Length of Sequence = 423 Nucleotides)

TITITITICCA TITAATAAGI ACTITATIGA TATTATATCA CACAGCACTI TACAGIATAC TCAAAGATAG CCIAAATTAT
GAATTAAACA TGCAAATATI TNCTITICCA AAATGIGGAC AAAATGICIT TIAGAGIGCI TITIGAACACI AGCCITAGCI
ACTAAGCATI CATGGGITTG ATCTITCTTG CGACATGACT TIAAGITAAGI TAACAAAAAA TGTAGCTGTA GACAGTAATT
GITTGATAAA TATGANCAGI TITIAAAATGG CACTGAATTI ACATCTITAA TCATTITTAAT AGGGCCATCC ACAGCCTCTC
TTGTGTCTCT AATTCTCAAC CTCCGGGGTC TITIAAAGGGC TGGTAAAGGC TCAGAAAGTG NCCAGCTCCA TGTGGGGTCT
CTGTAAGNNG TCTATGTCTT CAT

SEO ID NO:2366: (Length of Sequence = 294 Nucleotides)

SEO ID NO:2367: (Length of Sequence = 393 Nucleotides)

ACGGACAGAG CGAAGGGGAG AGGATGGTAG TGTCTGACTT CCACGTTTTC GTCAGGGATG TGTTGCAGCA TGTGGATTCC
ATGCAGAAAG ACTACCCTGG GCTTCCTGTC TTCCTTCTGG GCCACTCCAT GGGAGGCGCC ATGGCCATCC TCACGGCCGC
AGAGAGGCCG GGCCACTTCG CCGGCATGGT ACTCATTTCG CCTCTGGTTC TTGCCAATCC TGAATCTGCA ACAACTTTCA
AGGTCCTTGC TGCGAAAGTG CTCAACCTTG TGCTGCCAAA CTTNTCCCTC GGGCCCATCG ACTCCAGCGT GCTCTCTCGG
AATAAGGACA GAGGTCGACA TTTATAACTC AGACCCCCTG ATCTNCCCGG GGCANGGGCT NAAGGTGTGC TTT

SEO ID NO:2368: (Length of Sequence = 187 Nucleotides)

GATCTIGAAG TIAAACCAGI GITAGAAGIT TIGGIGGGGA AGACAATINA GCAGICTCIT CIGGANGIAA IGGAAGAAGA AGAGCIGGCI AACCIGCGGG CCAGICAGCG IGAGIATGAA GAACIACGGA ATAGIGAACG TCCIGAAGIT CAACGACIIG NAGAGCAAGA NAGGCGACAC CCAGAAG

SEQ ID NO:2369: (Length of Sequence = 341 Nucleotides)

GTATCTTAG TAGAGGGGG GTTCCACCAT GTTGGCCAGG CTGGTCTCGT ACTCCTGACC TCAGGTGATC ACCTGCCTCC

TCGGCCTCCC AAAATGCTGG GATTACAAGC GTGAGCCACC GCGCCTGGCA CCATCAGTTT TTGATCCTGA TACTTGTCTG

TCCTCTTGGT TCTCCTCATC CCTAATTTAA CCTTGAACAC AAAATTCAAC AGGTTTTGGC ATATAGAATA AAGATTATCA

GGCAAAGGCG CACTCTTGAC CTAATGATAT ATCTACATTT CATTTCCTGA TCTATCAGCA ATATTTAATT TGTCTAGAAA

TGATGAGAAG TTTAGAGGAG G

SEQ ID NO:2370: (Length of Sequence = 337 Nucleotides)

AGATCAAGAT CTCTCCCAAA ATGCCAGTAT GCAAAGGACA CTTGGGGCAG CCTCTCAACA TTTTCTGCCT GACTGATATG
CAGCTGATTT GTGGGATCTG TGCTACTCGT GGGGAGCACA CCAAACATGT CTTCTGTTCT ATTGAAGATG CCTATGCTCA
GGAAAGGGAT GCCTTTGAGT CCCTCTTCCA GAGCTTTGAG ACCTGGCGTC GGGGAGATGC TCTTTCTCGC TTGGATACCT
TGGAAACTAG TAAGAGGAAA TCCCTACAGT TACTNGACTA AAGATTCAGA TAAAGTGAAG GAATTTTTTT GAGGAAGTTA
CAACACACAC TTGGATC

SEQ ID NO:2371: (Length of Sequence = 320 Nucleotides)

CGTGGCCGCA GAGGCAGCTG AGCATGAGGG ATGGAGCGTG CTGCTGTCCT GCAGGTGCCG TTAGCCCTGT TTTGCACTGG
TGGATTGATC TGCTCAGGCG CACAGGGAGA TGGCACAGCA GGACCCGCCG CCCAGCCTCG CTGAGGGCAT GCTCCCGCCT
CACCTCCAGA GGCTGTTGGG CGGAAGCGAG AGCTGCAGCA GTTGGGGCCA GCNTGGGACT GGAGGCCCAG GTGAATCTTG
TGGGGCAGGG GACGGAGCTN AGGCTGTCCG GCCCGGGCCC TTCCCACCCA AAGGCCCTAG AACCCTAGGC CTTCAATCCT

SEQ ID NO:2372: (Length of Sequence = 326 Nucleotides)

AGGCCTGGCA TGCGGCGAAA AGTTCCTGGA GAAGGCCTCC CCCTCCCCAA AACACCCGAG AAACGTGGGG ACCTCATTAT
TGAGTTTGAA GTGATCTTCC CCGAAAGGAT TCCCCAGACA TCAAGAACCG TACTTGAGCA GGTTCTTCCA ATATAGCTAT
CTGAGCTCCC CAAGGACTGA CCAGGGACCT TTCCAGAGCT CAAGGATTTC TGGACCTTTC TACCAGTTGT GGACCATGAG
AGGGTGGGAG GGCCCAGGGA GGGCTTTCGT ACTNCTGAAT GTTTTNCAGA GCATATATTA CAATCTTTCA AAGTCGCACA
CTAGGA

SEQ ID NO:2373: (Length of Sequence = 361 Nucleotides)

AGCAGAGCTG AGGGAAGGCG TAGGATGGCT CCAGCTTCCG GTCAGTGGCT ACATGGTCAG TTCCATGATG GCGTTGACGA
TGTCACTGTG GTTGINICTC AGAGCCCGCA CGGCCTTGGC CCTGGACACA TTGGCCTGCG CCATCACCAG CTCAATGTCA
CGCAGTTCCA GCCCGCCTC GTCCACCTCT TCCTCCTCCT CCTCTTCCTC TTCCTTGCAC TCCAGCCTCA CCCGGGGCCT
GGGTGCTGAC TCAGGGACCA AGGCTGAGGG CTCTGAGGGA ACCTTAAACT TCTCAGCTGC GGCTTTGTGC ACTTGCTGGG
ACAAGGTCCT CAATCTTGGN CTCGCCAAAG ACCACATAAG T

SEQ ID NO:2374: (Length of Sequence = 281 Nucleotides)

TGACTCTAGT CTGGCACTTA TTGATGACAT TGAGAGGCTG AAATATGAAA TTNCAGAGGT GATGACAGAG ATCGACAATC
TAACTTCCGT AGAGGAGAGC AAAACGACTC AGAGGNACAA ACAGATAGCC ATGGGAAGAA AGAAATTCAA CATGGNTCCC
AAAAAGGGAA TTCAGTTTCT AATAGAAAAT GACCTGCTAC AGAGTTCCCC AGAAGACGTC GCCCAGTTCC TTTATAAAGG
AGAAGGCCTA AATAAGACCG TCATTGGGGA CTACCTGNGG T

SEO ID NO:2375: (Length of Sequence = 391 Nucleotides)

ATGITTAGIG CITCCITCAG GAGCICIGGI AGGCCAGGIC TGGIGGIGAC AAAATCICIC AGCATTIGCI TGICIGIAAA GGATTITATI TCICCITCAC TIATGAAGCI CAGITTIGGI GGATATGAAA TICIGGGIIG AAAATTCITI TCITTAAGAA TGITGAATAT TGGCCCCCAC TCICTICIGG CITGIACAGI TICIGCIGAA AGATCIGCIG TIAGICIGAT GGGCTICCCI TIGIGAGIAA CCCGACCITI CICICIGGCI GCCCITAACA TTITINCCIT CATITCAACI TIGGIGAATC TGACAATTGI GIATCITGGA GITGCIGITC TCGAGGAGGC AACCITTGIG GGCGITCICI GIAATTICCC CGAATTIGAA A

SEQ ID NO:2376: (Length of Sequence = 324 Nucleotides)

CCAGCCCTCC CTCAGCTGGG AACACAGCCA GGTGCCCTCA GACCCCTGGN TCTGCACAAG GGGGGCCTGC CCCCTCGCCC CAGCTATATA CACGACAGCC CATCCTGCTG GCCGTGGACA AAAGCTGGGA GCTCCTGTGC CCAGTAGAGA GCCCCTACAG TCCACCAGCT GCGCGGCCGG GTCCAGGGGC CCACTGTGGT GCCAGCNAGT TTNTCAAAAC CNAGGGCCCA GCCCCAGCTG GCNCCTNGCC AAGCCCCAGG CCTGTTTGCT GGGATGGAGC CTCCACACTG AGGCTGGTAA AAGCTTGAAC TCAACAGCAG CAAT

SEO ID NO:2377: (Length of Sequence = 357 Nucleotides)

GTTTATGTT TTATTTATGT ATTITAACTG ACTTATTTGT GTATCCCACT AGAACAATAC ATTCACAATA TACTTGCAGA ACTGTGCCTG GTGCGTCATG GGAGCAGAGA ACTTGTCCAG TGAATAGTTG TTGAAGAAAG GAGTAAAATC TCCCCCAAAC CCTAAAGGCA TCCTTTTCGT AGTGTGTGC CCATAGGTAT GGCTGCTGAG CACCAGGGCT GCTCACCATG CTCCCAAGAA GCAGAGTCAG GGAGGCAGAC AGCAGGGTTT ATTAAGGTGC ACACCCATGT CTGAGCCCCA GCTCTCTCCG NCTTCTGTGG GGAGGAAGCC CTCCGGTCTT TCCGAGGAAC CTTCAAA

SEO ID NO:2378: (Length of Sequence = 454 Mucleotides)

GACGGGTCTT TCAATAGCAA GTTTTCACTT CATCGACAAC ATCACGAAGG TEGTAACAAA CAAATGCTTT ATCAGGCTGG
ACTTCATTTC AAACTCCCCC AAAGCACAGA TCCATTACGC ACATTTAAAG ATACCATCTA CCTTACTCAG GTGATGCAGG
CCCAGTGTGT CAAAACAGAA ACTGAATTCT ACCGCCGTAG TCGCAGCGAG ATAGTGGATC AGCAAGGGCA CACGATGGGG
GCACTTTATT GGCAGTTGAA TGACATCTGG CAAGCTCCTT CCTGGGGCTT CTCTTGAGTA CGGAGGGAAA GTGGAAAATG
CTTCATTACT TTGCTCAGAA TTTCTTTGCT CCACTGTTGC CAGTAGGCTT TTGAGGGATG AAAACACGGT CTATATCTAT
GGGTGTGTCA GATCTTCACT CGGATTATTC GATGACACTC AGTGTGAGGA GTCC

SEO ID NO:2379: (Length of Sequence = 224 Nucleotides)

GGAAGAGACC TCACAGGINA TTAAANGIGT ATTTINIGGA CCIGGGCTTG GCTGGAATGC TCAGGGGTCC TGAAGATCCT
ATTATAGCTT CCITCIGITG AACCATTAAG AAAAGATGGC GANAGTCAAC ATAACTAGAG ACCICATCCG TAGNAGATCA
AGGAGCGGGG TGCCCTTAGC TTINAGCGGC GCTACCATGT CACIGINCCC TTITATCCGGC GGCT

SEO ID NO:2380: (Length of Sequence = 274 Nucleotides)

AGGITIGAAA TATCTTTTIG CAATAGATAA TCTTATTTAC ATTAATACAG AATCATTTTA CATTCCTAAA TCAGACACTA
ATAGATGCTT TATTTTAGTG AATTATAAAG GAAAACAAAA AGGAAACTGT TGAGAAGTGT TCTTCATTAA CCNGTCTAAC
GNCAGCCCGA AGATCCNGNA ACACATGGAA ACTGCGNCAT GCTNCCNGCA GAGGCTGGGG AATGGGGGTT CTGCTCTCAC
TGAATGGTGG GGAACCTTCA ACTGCTTAGC CTGT

SEO ID NO:2381: (Length of Sequence = 312 Mucleotides)

GCACAAACAG TTTTATTTGA TGANCCACAG TGACTAACAG GNTCAGAAGA CAGTGCAGAT ATTCTGAAGA AGGCACTGNG
GGAGGTAAGG GGGTATCACA GCAGGCAGCC TCCTCTENTT CINTCCCAGT TCACAGATGA GTTCCAGGCA GGAAGTCTCT
GCAGGTCACC CACGGCGGCC TCAGAGGGAC AATTINTTCC CTTCTAGAAG CCTNTTCCAG TGTTCACTGG ATGNTTTGAG
GACAGNTCTG GGCAGAGGAG GTGACTCTGT GAAAGATGCT ATCTTAAGAT GGGGAGACTA GGCTGTGAGG AG

SEO ID NO:2382: (Length of Sequence = 402 Nucleotides)

CTTAAACTAA CTITGAAGCA AGTAATGICA ACTITGAGCA CTITGITGAG TITTGAAAAA TCTTATTIGI TGCTGCACAG
GITAATAAAT TATCAATTIG TAATICAGCA TGITGGICAG AGACACGGIC ACTGATICAC ACCCAGTCCC TGCCACAGAC
CGICTCAGAC ACGCACAGIG GGCCTGCTGC ATGATTCACA CCCAGTCCCT GCCACAGACC GTCTCAGACA CGCACAGTGG
GCCTGCTGCA TGCGTGITAC CTGGCTTTTG GCTCCACGCT CACTCATAGC CATGTCCACA TGGGGGGCTT GCACACAGGA
TCACTCACAT ATGTACATGT ACCCACCACA AACGTGCAAA GCTCCTTGCA CACATGCATG CACACAAACG TGGTACACAA
GT

SEQ ID NO:2383: (Length of Sequence = 406 Mucleotides)

GACCCITTIC ACITAGCCCI CITEGGITTE CAACAIGCIT TCTCTCCAC CITCTCATTE AATGAGAAAA AACAGCCCAG
CCAUTITTE CAACAGCAA AGCACCAGAG TEATHATEGC TITECTCATC TCACTAGTEA CITCACAGTAA CITCACAGTAA
TGTAGGCAGT CCAGGCATTA TTATTTTCAT TITACAGATG ATGCAACTGA GGCTCAGTGT GGTGAAACAT TTGGCTCATA
GCCACCAGGC TGATAAGCAT CAGGGACTTG GGACCTAGGN CITCACATTT CAAGTCAGCT GTATCTGTCC CCAAGCCCCA

CCAGACTTCA TGTGAAGGTG GCTGCTTCTG GGGTGATGGT GCCTGGAGAG GCAGACTTTG AGGCTGCCAT GCTCTTATTT TCAGAT

SEQ ID NO:2384: (Length of Sequence = 165 Nucleotides)

TTAAGACAAT AATGAAAGAT TCTGTACAAA GTTACCAAGT CTACAGGCTG AGCGAGCCAA GGGTAAGTGG GGCCTGATCC TTGTGGACGA ATGTNCCCGG GAGAGCTGGC CTCACCTGGG GGAGGCACGT TGAAAAAGTA CACATTTACA GGGCTCGGGA AAGGC

SEQ ID NO:2385: (Length of Sequence = 297 Nucleotides)

GGTTTINATT CATTCTCTTC TATTAACCTC TCTAAAGGAA ATTGGGCACC TGTAATCCCA GCACTTTGGG AGGCTGAGGT
GGGTGGGTCA CTTINAGGTC AGGAGTTCAA GACCAACCTG GCCAGCATGG TGAAAACCCA TCTCTACTAA AAATACAAAA
MITAGCCAGG CTGGTGGTGT TCGCCTGTAA TCCCAGCTAC TCAGGAGGCT GAGGCAGGAG AATTGTTTGA ACCTGGGAGG
CGGNGGTTGC AGTGAGCTGA GATCGTGCCA CTGCATTCCA GCCCAGGGTG ACAGAGT

SEO ID NO:2386: (Length of Sequence = 290 Nucleotides)

AAAAAATAAA GIGAATITAT TGGITCATGI AACTGGAAAG TCTCATGAAA ATGICAGCTI CAGGAGAAGC TTGACCCAGC
AGCITCATGA TGIATGGAAA TACCTGGGIT TITTGITTCT NCTCTGCTAC TGIGGIATCA GCTTTATTCC AAGICTGGCT
TCCTTTGITG TTGCAAAATG CTTTGICAGA AGAAGCCTGG GTCCATCTGT TAGGNITAAG TTTACTCTGT ATGCTGTAGT
AGTGGCTATG ACAAGATTAG GAAGTGTATT TTCTCCTCCC ATATTAAAAG

SEO ID NO:2387: (Length of Sequence = 356 Nucleotides)

GTCATCTGTA TTGTCACATG AAATGCACAT CCAAAACGGG TGACTTGGAA ACGACCTATT AGGTCACACG GAGTCCGGCC CCTGGGGGCA AAGCCTCATC GATGCCCACG GGCGGTGGCC AGCACTTTCC TTGGGCTGTG GCGTGTGCAC CCGGCCTCCC CAGCGGAGAG TCAGCTCACA CCCCAAGGCC TTTAGCTCTC TGGCAGCAGC TCCCAAAACG CACTTGAGGA ACCAATAATT CCTTGGGGGT TAATAGCTGT TCCCCAAGAA AAGCGTTCTG TGGGTCAAAT AAGTTTAGGA AAACATGGGT TAAAAGAAGGT TTAGGCAAGA AGCTTTTCTA TAGGGCTTTG TCAGAG

SEO ID NO:2388: (Length of Sequence = 226 Nucleotides)

ATTATTEGTA TAAAAACTTA AGACGGCATT AGAATTCTTA AGAAAAGGTG TAAAATTTAA AAAGATGTGC AAACAACAAA GAATGCCCGA CCCTGAACCA GACCTAAAGC ACCTTCCANT TCCTCCACAC ATCATGCCCC AACACCATCC AGCCCAATCG GACACCAGGA CAGTGAGGGA CGGGTGGCTG TTCAGTGGGC AACAGATCTG GAAGGAAAGA TTTTCA

SEO ID NO:2389: (Length of Sequence = 250 Nucleotides)

CCCAGCTAGG CCITGGRIATG GCTNCAGTGA GGAGAAATCC CGGGAACTGT ATTGACACAA AGATTCTNAT TGCACTTGTA
TTTTTNTATT AAAGTTTGCA TGGTTTCTAA TAAAGGATTC AAACATAAGT TTGTAGTGAA ATGGCCTGCN AGATTCCAAG
GGCTTCTCTN GAAGGGGGAT TGNGCTGCAN TGTAGATTTN CCTCTGAAGG AGGCTGGCCC CAAACTTGGN CCTCCTCATG
ACCCCCTCCT

SEO ID NO:2390: (Length of Sequence = 371 Nucleotides)

CCTTTTCIG GAGAACGGG TCTCGCTATA TTGCCCAGGC AGGTCTCGAA CTCCTGGGCT CAAGCTATCC TCCCGCCTCT NAGCCTCCGT TTCCAGAAGG TCACCAAGTA ATATCTGCNT TTCATCAGTT GCAGTTAAGA TTTTNNTTTC TTGAAATACT GGTTTTCAAA CAGATCAGAA TTACCTGGGG AGCTTGTTTA AAATATAAAT GCCCCAAGGC CAGCTCCAGG ACATTCTGAC

GAATTITIT ATTIAGGIGG ATGCATTITI NGICTGITTA CIGCTCTTCT CAGCTTTATT CAATAAACTT GCATTITAAG
GGTTGTATTG GCAATTITAA CITAAAATGT GCATCATGAT GGAAGGIGCA GACCTTTIT

SEO ID NO:2398: (Length of Sequence = 421 Nucleotides)

SEO ID NO:2399: (Length of Sequence = 392 Nucleotides)

GATAAGCTTG ATATCGAAAG TNCCACAATG GGTCAGCTGT ACCAGGAACA CCATGAAGAA GACTTCTTTC TCTACATTGC
CTACAGTGAC GAAAGTGTCT ACGGTCTGIN AAGCTGCTGC CCCTGAGCTG GAGGGGGGTC TCATTCTACA AAGAGAGAGG
TGGCCCCCCT TTCTTGACCT CCTCCTCCTT CAAGCTCAAA CACCACCTCC CTTATTCAGG ACCGGCACTT CTTAATGTTT
GTGGCTTTCT CTCCAGCCTC TCTTAGGAGG GGTAATGGTG GAGTTGGCAT CTTGTAACTC TCCTTTCTCCC
TTTCTCTGCC CGNCTTTCCC ATCCTGCTGT AGACTTCTTG ATTGTCAGTC TGTGGTCACA TCCAGTGGAT TG

SEQ ID NO:2400: (Length of Sequence = 366 Nucleotides)

CTGGGGAAGG ACTGGCACAA GTTCTGCCTN AAGTGCGAGC GCTGCAGCAA GACGCTGACG CCCGGGGCC ACGCCGAGCA TGACGGGAAG CCGTTCTGCC ACAAGCCGTG CTACGCCACC CTGTTCGGAC CCAAAGGCGT GAACATCGGG GGCGCGGGCT CCTACATCTA CGAGAAGCCC CTGGAGGAGG GGCCGCAGGT CACCGGCCCC ATCGAGGTCC CCGCGGCCCG AGCAGAGGAG CGGAAGGCGA GCNGCCCCCC GAAGGCCNCA GCAGAGCCTC CAGTGTCACC ACTTTCACCG GGGAGCCCAA CACGTGCCCG CGCTGCAGCA AAGAAGGTGT ACTTCGCTTG AGAAGGTGAC GTCTCT

<u>SEO ID NO:2401:</u> (Length of Sequence = 385 Nucleotides)

SEO ID NO:2402: (Length of Sequence = 392 Nucleotides)

AAAGAACTIG GTATCICTAT TAAAGTACAT GANCCICCAA GGAAAATAGA GCGATTIACT CITCICCAAT CAGIGCATAT
TTACAAGAAG CACAGAGTIC AGTATGAAAT GAGAACACTI TACAGATGIT TAGAGTIAGA ACATCTAACT GGAAGCACAG
CAGATGICTA CITGGAATAT ATICAGCGAA ACTTACCIGA AGGGGTTGCC ATGGAAGTAA CAAAGACACA ATITAGAACAG
TTACCAGAAC ACATCAAGGA GCCAATCTGG GAAACACTAT CAGAAGAAAA AGAAGAAAGC AAGTCATAAA GCCTTCAGGG
AGGCCATTIT TGCCTAAATT TTGAAATGAG GGTGGGCCCAG ATGAGTATGT TTAAGTGGAG AGTGCTTTCC AG

SED ID NO:2403: (Length of Sequence > 179 Nucleotides)

SEO ID NO:2404: (Length of Sequence = 399 Nucleotides)

TTCCCAAAGT GGITGIAACA TITTACACTC CTACTAACAG TGCATGGGAA GCCAGITTCI CTATATCCIC TCCAACATTT GGIGCIGICA ATCITTAAA ATTTIAGCCA TITTIGIGGI TGIATAGTGI TATCTCATTG CAGITTIAAT TTGCCGATCC CTGAATGIGT GTAGGIGTGI ATATGIATTA TATAATATAT ATATNATNCI TTCACTTATT TTGAAGTAAT TTCAAAGTTT CCAGAATAAT ATCAAGAACT CCTGTACTCC CTTCGCCAGA TTCTCCAATT GTAATGITTT ATTGCATATG CTCCATTGCC CATTCTCCC TCTACTTATA GCTTGCATTA GTGTTTTCCT GGAACCNTTA GAGATGAAGG TGGAAAAAAG GATGCGGGT

SEO ID NO:2405: (Length of Sequence = 404 Nucleotides)

GGAACAGAGT GACCTGACCA CCCTAACATC AGCTGCATAC CAGCAGAGCC TGACTGTTCA CACAGGAACT CATCTCCTCA
GCATGCAGG GAGCCCTGGA GGACACAATC GCCCAGGCAC CCTCATGGCA GCTGACAGAG CCAAACAAAT GTTTGGACCC
CAAGTGCTTA CGACCCGGCA CTACGTGGGC TCAGCAGCTG CTTTTGCAGG GACACCAGAG CATGGACAAT TCCAAGGCAG
TCCTGGTGGT GCCTATGGGA CTGCTCAGCC CCCACCTCAC TATGGGCCCA CACAGCCAGC TTATAGTCCT AGTCAGCAGC
TCAGAGCTCC TTCGGCATTC CCTGCAGTGC AGTTACCTAT CTTCAGCCAC AGCCACAGGC CTATTGCTGT GCATGGGCCA
TTTTT

SEQ ID NO:2406: (Length of Sequence = 280 Nucleotides)

AAGAGAGAAC ATTITATITG TCTATAATTA GGGTAAACAG TTGGGTAAAA YCTTACTAAA AGAAAGITAA GGITGTCTTA
ACACAAGATA TATAATGACA TAAATYAGIT AATTAAATIT YAATTAAAAM CAGCIGCTIT GGAAATCCAA CATGTATACT
TCAAAATAAT TTACCTAAAT AACTTATGAA AATGGATGIT ATTGTACAAC TCATCTCTCC TTATAAAAGG NGAACAAAGG
ACATAGGAAA GCTGAAAAGA AGGCTAGATG AAGATACAGG

SEO ID NO:2407: (Length of Sequence = 350 Nucleotides)

TOCAAGGGCA ATATAAATTA CAGTATGCAA AACATACTGA CTGGCTGAGG TAAAACGCAC TGCTCCTGCC TCACGTCACC
ATGAGGGGAA ACACACATAT GCTTTTAAAA ACATCTGGCT TATAAAAAAAA CATCCCCTAG AAAGGCCTCC AGAGAGGGGC
TGTGAGGCTC ACCCTCTGCC GCGCTCAGGA GGACCCGCCG GCTCAGCCCT GGCCCCTCCA CTGCAGCCAT GGGTGGCGCC
TCCCCCTACT GCCTGCCCAG GGCTCTGTCC AGGTTGCTCT TGATGGTGTC GAGGAAGTCC GTGGTGTTCA GGAAGTGCTC
GTTCAGCCTTC ACATTGCTGA GGCCGTGAAT

SEO ID NO:2408: (Length of Sequence = 239 Nucleotides)

ATMENITIGE GETCECNAGA AATEGATETE CEGAAGAAGA AGAAGAAAAA AAATCAGCAG CIGAAAGANC CAGAGGCAGC AGGGCCTGTG GGGACAGAGC CCACAGTGGA GACACTGGAG CCTCTINGNAG TCCTGTINCCC GTCCACCACC AAGAAGAAGA AGAAGCCCAA AGGGAAAGAA ACCTTCGAGC CAGAAGACAA GACAGTGAAG CAGGAACAGA TTAACACTGA GCCTCTAGA

SEO ID NO:2409: (Length of Sequence = 331 Nucleotides)

TCTCTTCAAG AATTTCAGAC CAATCGACCG TCCTGTCTCT TTAAGGCTTA GGAAGAGCAG TGTGGCTGCC CCTTTAAGGA GGCGTTGCAA CAARCCATAT TGGACAGACG ATGGGGGCCGA (XXATCGGGA CCCGALGGGC CTCTGACTCC AGCAATATAG CGAATCAGCG GCTTTCGGGA ATACATTTTT CGGAAAAAGA CTTCTTCCTC GGTTTTCTGC TCTGCACACG TTGAAATTTT CCCCAGTTTT TCCTGCAGAT CGGGAGTCGA GCAATGCCTA CCCCCGCCTC CCGCACCAGT TGGGCGCTCC CGGATGATGC CCTACCCCTT T

SEO ID NO:2410: (Length of Sequence = 135 Nucleotides)

CTGCAGGACT TGCGAAGAGC GTGCATTCCC AGTGGGCGAA CGGGAATTCG AACGGAGAGA GGGTTATCTT GTGGGGGGGCT ACCGGTGGAG AGCAAGGGGG CCCCCAGGGGT TGGATCGGTG AAATTNAGGT CGCCC

SEO ID NO:2411: (Length of Sequence = 330 Nucleotides)

ATGCTGCTCG GITCTCTTGT CCCCCCAACT TTACCGCGAA GCCCCAGCCT CAGAGTCCCC TCGTTTCTCC TTGGAGGCGC
TGACGGGTCC AGATACGGAG CTGTGGCTTA TTCAGGCCCC TGCAGACTTT GCCCCAGAAT GCTTCAATGG GCGGCATGTG
CCTCTNTCTG GCTCCCAGAT CGTCAAGGGC AAATTGGCAG GCAAGCGGCA CCGCTATCGG AGTCCTCAGC AGCTGTCCCC
AAGCTGGAGA AGCGACCCTG CTGGCCCCCT CAANGGAGGC AGGAGGTGGA CTCACCTGTG CCTCAGCCCC CCAGGGCACC
CTAAGGATCC

SEO ID NO:2412: (Length of Sequence = 583 Nucleotides)

TECACCETE CACCAGETEC COSTETEGAT TETNACAGNN ACTEGETINA TGAAGETIAC CACCTACCEN GTECACETEG CONAGCAGCA GGACGTECAC CTEACTETINA CEGAGTCTCG GCAGCATGAG CTCTCGCCAG ACTCGAACTT GCCCGTGCAG CTCCTCACCA TCCGTGTGGC CAGCACCAAC CCTGCTGTGC AGGCCTTTGA CATCTGGCTG AACTCCACTG AGTACGGGGA GCTCTGCGAG AAGCTCCGGG CACCCATCCG CAGGGCAGC CATGTGGTCA TCCACCAGAG CCTGGGCGAC CTNTINNIGG AGACATTTGC CTCCCTGGTA GAGGTCAACC CGGCCTACTC AGTGCCCAGC AGCCAGGAGC TGGAGGCCTG CATAGGCTTG CATGCAGAACA CGTGCCAACG TGAAGNTGGT GAAGACCTGC CAGGAGTCAG CCACAGGGGA GTTCCAGCAG TNTTAATTNC CGCCCCATGT TGGTGGCTTA ACTTGATNGG GAAAGTGGNT TNGNCAAGCG GCAAGACCCC CTTGGGCNCTT NAAACTTGNT TGGCCAAACGG GGTNCCTGCA TGG

SEO ID NO:2413: (Length of Sequence = 203 Nucleotides)

TOSTOCTOCC ACCOCCTAGO CATGCAGNEG TGAATNEGGG AACCOAGGNN GGGGGCTGAG AAGCTOCAGG CCACCTTNAG GGAATCCACG AGGGTCTTTC TACCAGGAAG AAGTGCCGCA GCTGCGTGGC CGCCGAGACC ACGCGGGAGG TGATCTGGTG GGACAAACGT TCCGTCTGCT CCCGAGTCAG GAGATCGAGT CTC

SEO ID NO:2414: (Length of Sequence = 92 Nucleotides)

AAGGGCAGG ATGGGCCTGG GAAGTCCAAC CCCACGCATT TGGGCTCAGC CTTGGACATG GAGGCCTGAC AGCTGTTGTC

SEO ID NO:2415: (Length of Sequence = 401 Nucleotides)

CTTTTCCCTT CIGIGENCCA AATGCANCAT CTINATACAC GITGCTTAAC CTAGAANCGT GGCTCCACCG TGAATTCTAA
TTGGTCCGIG CTATCGAGGC ACTGTCCCCT TAACTGGTCT CGCTCCAGIG GCCCCNACTG CTTTTCTTCC TCTTCCAGNA
ATGGCTCTTC GGGCCCAGAG TTCGAATCTC GCGATCGGGA TGGGGACGGA GTACCGGCCT GGGGTGTCCC AGAGCCCGGA
CTGAGCTGGG GAGTCAAGAC CTCGGGGGAT GAGGCCTGAG CAAGTCGGAG TCGTAGGTCC AGTTCTTCCC
TGTCTCCAAT CIGTTGGGTT CTTGGGGTTC TTCGTCTTCC AGCGGGGGG AGCTGCTGGT GGAAGAGTCC TCCCCGGAIC

SEO ID NO:2416: (Length of Sequence = 245 Nucleotides)

ATGIAATACA GIGIAGAAAG CGATCATGIC ATAAGCAATG ATTCTGIACA ATCATNCNGC AGAAAATTAG TITTGGAGAA TICITGGIAA TIGAAGACCA GCAGAGCACC CCTCCCCACC CGCCCCCTAA AAGIGCTTAC AATTTACAGG GATYCTITTC TITTTCAAAG ACCCAAAGAY ACGIGGICAG AAAAMAAAAG CITGAAGICT CAATGCCTAA TGICGIGCAC ATTKNACAGG GACGC

SEO ID NO:2417: (Length of Sequence = 384 Mucleotides)

GGITTIGCAA GATGATGGAA CATCCCATAA GCCCAGGIGI GCAGCIAACC TITAGAAGCT GGAAAAGGCA AGGAAACATA
TICIGIAGAG CCICCAGAAG GAACACACGI CIGCACACAC TITGITTITA GCICAGIGAA ACIGATITIG GACIACIGAC
CITCAGAACT GIAAGATAAA TICCIGITGI TITACGITTG TGGIGITATA GAAGITACAG AAATGAATAT ACITACCGIA
GTTIAGAGAG AGATGGGAGG ATACTITTIT TICICCCTIC TITTIGAAGG GAGGIAGGIC TCCITAACIC CAGAGGAAAG
ACTIGICITT CITCATATAG GGGCCCTTIG ATTCITAATT CATGGGAGT GITTAGGAGA TIGA

SEO ID NO:2418: (Length of Sequence = 1645 Nucleotides)

GTGATGGCTG CCTTGAGGGG GACCATCATG TOGGAGACCG CATTGGTGCA GGTCTCACCC CACAGCCCAT GCCCAGCCTC CIGCAGACTC AGGICATCCA GCIGGICGAT GGCICTITGC ATACCIGGIG CCITCTCCIC TCGGGCITGG CAGGCITCTC TEGEGGCTTC TCAGATGACT CTTTTGCCTT CTTCTCTGTC TTGGCTAACT CCTTGGCCAG CTCTGAACGT GCCTCCTTGG CTCCCTCTTC TACCACCTCC TCCCGTTTGG CCAACTTGCT CACGGCCGTC TTGGTAGTGG CTTTGAGGCT CTCCTTGCTA TCAGCCCGCT GTTTGATTTT GCTGGGCTTG AGGTTGGTAG GCACAGCCCC AGAAGCCAGG NCCTTCTGCG TGGCCACAGG GTAACGCAGG AAGTCCAGAT GCCGAAGCTT TTCTAGGCCC TCCAAGATCT TGTTTTGGGG AGCATTTCCT GGAAAAAGCA CACGCACAAT CITCICAGIG GGATIGGCIG GIAGCCAGAC CACCAGAGCA GIGAIAGAGG TAAGGIAGGG CACGGAGAIC TCAGCCTCCT TCCCATTGGG CAGCACGATG CCTGINITGG CITTACTATT GCCTGCCCAC TITTGCATGA GGAACTGCAT CTCCTTGCTG TCCTTGACAG GGTTGAGGAC ATACATGTCC AGCCGGCCCA CACCCATTTT GTGGAAGAGG GTCAGTGGCT CAATGGTATT GCTGACCACA CGATATAGAG GCTCAGCCTG GATGCCCAGG CGGTTTAAGT GCTGCAGAGT GAGGCAGGCC TCCTCAATGC TACGCTTGGC TTTCCGGGAG GCATCAGGAA GCCGCAGCTT CTCAGGCACG TTGAAAAAGA CAACTCCAAG CTCAGGANAG ATAAGGITCT TCACCCAGTC GCTGTAACTG CTAGAGCCCT GGNACTGCTC CTCCTCTAGC TCTGCCACTT TECECTECAG TAGTCCATTG ATECCTESCA GETTETCTEC CCCAATGTET GINAGTAGCA CCGAGTCAAT GCCGTCCAAG TNCCGIACCA GCITCCAAAA ACAGGACTIG CGATCAGAGC CACCATCCAC CAGGATGITG AAACCATTGA CAGCAAAGAG GGCAGAGTCC CCACGACCAC CTGGGAAGAT GTAGCAACAA GGCTTGGAGA GCCTTGAGGAA GCCCCCTGAG GTGGGGGGCT CTAGTAGGTC AAATGGGGAT GGCACGTCCA CAGTCTCAGA GACATACTCG GAGAACTCAG CCACGCCGTC CATGGTGGGC AGAGTGGGCT CAGGGTTTAG CCGGAGGTGC AGGGTCTCTT GGGAACTGGA TAATCCCAGG TGGCTCCAAT CACCTTCCCC TAAGCAGGAC ACGGTAAGGA AGGCCTGTAT CCCAGGGTCT CTATTGCTGA GCAATTGGGA AATCTCGGGG TTGTGAAGGA CCTGGGCAAA GITTTCATAT GAGTAGGTGC CACTCTGTAG GATGAGGTCT CCCCCAGGCT CTAAACTTTG CCCACTCAAG ATTAGTAGTT TATAAGCTGA TGAGCTGCTA AGAAGATGAT GAACCTCAGA GCTGATGCTG TCTGCACTGG GATTTACCAG GATGATGGIC TCTAGGATCT CACTCTGGTG GCAAAGGGTC CTCTG

SEO ID NO:2419: (Length of Sequence = 837 Nucleotides)

GEAAGGATGA GAAACAGATT TINTECTCACT TCATGGGCTG GCCTGGAATT GACGATGGTG CAAACCCAAA TINATCCTGAT
GTAATTINATG AAGATTATGG AACTGCAGG AATGACATCG GGGACACCAC GAACAGAAGT AATGAAATCC CTTCCACAGA
CGTCACTGAT AAAACCGGTC GGGAACATCT CTCGGTCTAT GCTGTGGTGG TGATTGCNCC TGTGGTGGGA TTTTCCCTTT
TGGTAATGCT GTTTCTINCTT AAGTTGGCAA GACACTCCAAA GTTTGGCATG AAAAGGTTTTG TTTTTGTTTAA TAAGATCCCAA
CTCGGATGGGT AGCTGAAATA AAGGAAAAGA CAGAGAAAGG GGCTGTGGTG CTTGTTGGTT GATGCTGCCA TGTAAGCTGG
ACTCCTGGGA CTGCTGTTGG CTTATCCCGG GAAGTGCTGC TTTATCTGGGG TTTNCTGGTA GATGTGGGCG GTGTTTGGAG

GCTGTACTAT ATGAAGCCTG CATATACTGT GAGCTGTGAT TGGGGAACAC CAATGCAGAG GTAACTCTCA GGCAGCTAAG
CAGCACCTCA AGAAAACATG TTAAATTAAT GCTTCTNTTC TTACAGTAGT TCAAATACAA AACTGAAATG AAATCCCATT
GGATTGTACT TCTNINCTGA AAAGTGTGCT TTTTGACCCT ACTGGACATT TATTGACTTA ATTGCTTCTG TTTATTAAAA
TTGACCTGCA AAGTTAAAAA AAAATTAAAG TTGAGAACAG GTATAAGTGC ACACTGAATA GTCTAATCTA CATGTAACAC
ATATTTNNGT ATGATTTTCT ATACTCTAAT CAGCACT

SEO ID NO:2420: (Length of Sequence = 1843 Mucleotides)

GAAGCTCCGG CCCAGGTGGC CGCTGGCTGC TGAGCTCACG CCAAGGTGCG GCTGTGGTGG TGGTGGTGGC GGCTGCAGGC TTTGCTGCTG CTGGATGTTT GCTGGCTGCA GGTTCTGCTG CTGCATCTGT AAGTTTTGTG GCTGCACCTG CTGGGTCTGC ACCAGGTGAG GCTGGGTGGC CAGCCGGGTG CTGGGCAGGC CCTGGTAGCT CATCATCTGG GACAGGGCGC TGGCAGCAAG GCTACTGTGC AGCGGGCCTA CCATGCCATG CTGCAGGGAG GGGGCCTGTG TGCTCAGGGG GCCTGGTGCC ACACTCCCCC GCAGAGGGIT GTATTGGTTC GGCACCATGC CGCTCTGCAG CCGGGACAGC CACTCGCATT GACCATTCAA ACTGGTGGAC COGNICAÇÃO TGARATICAS OSCICITOGS CISCINGASC CUASGACESI SCIEGISCOA GASSICAÇÃO SCASSICEGA GGINCCCNAT GCCCAGGIGG GIGICGGGCA TYCCAGGCAG GIGGITGAGG GGCACGGACG GAGACTGCIG GAACGGGGAG GGCAGNAGTG GCGGCGAGGC CACGTCTGAC AGGTAGCCAT GGGGTGACTC CAGGGAGTCC ACGGCGAGA GCATGCCGGA GCTGTCCAGC AGGCAGNCCT TGCCGTCCTG GGACTTCTTC CTCCGTGCCT TGAGGTCCTT GGCCTCCTTG CTTCCACAGG CCAGGCCITT GCTGCTGGGC TTGCGGACCT TCTTGCCCTG CACGCCGGC TTGAGGCTGC CCAGGTAGCC GTTGGGCGAG CAGAGCENGG GCGACAGGGT GGGCGTGCCC CCCAGGGGGC TCCGTGCAGC TGCGGGGTGC GCACCAGGTT GTACTCGTCC AGCAGCCICA CGATGICGIG ATGCATGCNC TCCINTGCGA TGTCGCGGGG CAGGCGGTCC ATATGATCCG TGATGTCCCG GTTGGCAAAG TGGTCCAGCA GCACCTTGGC GGTCTCGTAG CTGCCCTCCC GGGCGGCCAG AAACAGGGGT GTCTCCTCCC TETTETTCTG CATATCTTTG TTAGCCCCGT TCTTCAGGAG CACAACTGCG GCATCCACAT TGTTCACNGC GCCGCCCAG TGCAGGGCGG ACTTGCCCAG GTNATCTACG GCGTTGACGT CGGCGTGTGA GTTGATGAGG TCCTCCAGCA TGCCCTCCAC GGCCAGGCG GCAGCCAGGN TCAGTGGCGT CGTGCCATCA TGCATGCGGG CATCCAGGTC TGTGGCTCGG TTCCGGATCA GGATCITGGA AGACACCTTG TGCGTCGGCA GACACAGCCG CATGCAGCGG GGTGCGGCCC ATGTTGTCCT GGATGTTGGC ATCTGCGCTG GCCTCCAGCA GGCGCTTGGC GGCATCAGAG CGTGAGTAGC GGGCGGCCAG GTGCAAGGCG GTCTCGCCCCG TNEGGTETGT CTGGTTGTGC AAGCTGGCGC CCTGGTAGAT GAAGTCGGAG ATGACGGCGG GCGGTCCTC CTCTTCCTCG CTGTTGCCCG TCTCCAGGCC GCCCCCGCTG CAGGAGGCGA TCATGAGCCG GGTGAAGCCA TCAGGCCCGC GGACATTGAC GTCCATGCAG TCGGCGTCAA CCTCACCCTG GGGCGGTGTG GGGGCCATGG CANACATGCG CAGGTCAGCG GCATCCAGGT GCTGCTGAGT CCACTGCCGG TGGTCTGTCT GGTCGTCCAG GTCAGGCAGA ACCACGGGCT CCTCGAACCG GAACTTCTTG GIC

SEO ID NO:2421: (Length of Sequence = 1452 Nucleotides)

GCTTTCTTCC	TACCCCATTC	CCCGCTTCCC	TCCTCCTCCC	CTGCAGCCTG	GTTAGGTGGA	TACCIGCCCT	GACGTGTGAG
GCAAGNTAAG	GCCTGGAGGG	TCAGATGGG	AGACCAGGIC	CCAAGGGAGC	AAGACCTCGC	GANGCARGCA	AGCCCCNGCC
							GGACGNAATG
CTAGCTGCCC	CTTTCCCGIN	CTGGGCACCC	CGAGINICCC	CCGACCCCGG	GTCCCAGGTA	TGCTCCCACC	TCCACCTGCC
CCACTCACCA	CCTCTGNTAG	TNCCAGACAC	CINCACGYCC	ACCIGGICCT	CINCCATOGC	CCACAAAAGG	GGGGGCACGA
GGGACGAGCT	TAGCTGAGCT	GGGAGGAGCA	GGGTGAGGGT	GGGCGACCCA	GGATTCCCCC	TCCCCTTCCC	AAATAAAGAT
GAGGGTACTA	AAGITGICIT	GGITTITATT	TIATTATTAT	TTTTTTTTT	TTCCAGTATA	CIACCITGIC	TTTTAAGAAA
GGGGATATTA	ААААААААА	AAAGACAAAA	GIGITITIAA	AAAAAAGCAA	CACCCACACC	TEGIGICIGI	ATATAGTCAG
CTTATCTCGT	GITCAATCGT	CIGATCICIA	CAGAGAGAAG	TGGAAAATGC	TGTATCAAGG	GIGGGCTTAG	CIGICCTIT
CCAATAAAGA							

20

25

30

5 WHAT IS CLAIMED IS:

1. A purified polynucleotide having a sequence designated as one of:

SEQ ID NO: 316 - 2421, except SEQ ID NOS 650, 1834, and 2073;

or having a sequence complementary thereto.

2. A purified polynucleotide having a sequence designated as one of:

SEQ ID NO: 316 - 2421, except SEQ ID NOS: 485, 650, 1834, 2073, 2092, and 2353;

- or complementary sequence thereto or, for those sequences over 150 nucletides long, a portion thereof at least 150 nucleotides in length.
 - 3. An isolated polynucleotide that includes a sequence designated as one of:
 - SEQ ID NO: 316 2421, except SEQ ID NOS: 485, 650, 1834, 2073, 2092, and 2353;

or complementary sequence thereto or, for those sequences over 150 nucleotides long, a portion thereof at least 150 nucleotides in length.

4. An isolated polynucleotide operably coding for a native human polypeptide or protein, which includes a region coding for the same amino acid sequence as a native human coding region corresponding to a sequence designated as one of:

SEQ ID NO: 316 - 2421.

- 5. The polynucleotide of Claim 4, wherein said SEQ ID NO is listed in Table 6 and is one of SEQ ID NOS: 316-2421.
- 6. The polynucleotide of Claim 4, wherein said SEQ ID NO is listed in Table 7 and is one of SEQ ID NOS: 316-2421.
- 7. The polynucleotide of Claim 4, wherein said SEQ ID NO is identified in Table 10 in a metabolic functional grouping and is one of SEQ ID NOS: 316-2421.

15

25

30

- 8. The polynucleotide of Claim 4, wherein said SEQ ID NO is identified in Table 10 in a structural functional grouping and is one of SEQ ID NOS: 316-2421.
- 9. The polynucleotide of Claim 4, wherein said SEQ ID NO is identified in Table 11 in a developmental control grouping and is one of SEQ ID NOS: 316-2421.
- 10. An isolated polynucleotide coding for a human protein or polypeptide, which includes a coding region corresponding to the EST identified as:

10 SEQ ID NO: 316 - 2421;

or a polynucleotide complementary thereto.

- 11. The polynucleotide of Claim 10, wherein the SEQ ID NO is 316-1000.
- 12. The polynucleotide of Claim 10, wherein the SEQ ID NO is 1001-1500.
- 13. The polynucleotide of Claim 10, wherein the SEQ ID NO is 1501-2000.
- 14. The polynucleotide of Claim 10, wherein the SEQ ID NO. is 2001-2421.
- 20 15. The polynucleotide of Claim 10, wherein said polynucleotide further includes the entire sequence designated as any one of SEQ ID NOS: 316-2421.
 - 16. An isolated polynucleotide comprising at least 150 bp of a sequence of Claim 10 and wherein said SEQ ID NO excludes NOS 485, 650, 1834, 2073, 2092, and 2353.
 - 17. An isolated polynucleotide sequence, which hybridizes to a sequence designated as any one of SEQ ID NOS 316-2421, except SEQ ID NOS 485, 650, 1834, 2073, 2092, and 2353, or to a sequence complementary thereto, under hybridization conditions sufficiently stringent to require at least 97% base pairing.
 - 18. A polynucleotide according to any one of Claims 4-17, in substantially purified form.
 - 19. A construct in isolated form comprising a vector and a polynucleotide according to any one of Claims 1-17.
- The construct according to Claim 19, further comprising a promoter operably linked to said polynucleotide.

10

- 21. A panel of at least 100 isolated polynucleotides having the sequences of Claim 3 or Claim 16.
- 22. An antisense oligonucleotide capable of blocking expression of any one of the polynucleotide-encoding sequences of Claim 10.
- 23. A triple helix probe capable of blocking expression of any one of the polynucleotide-encoding sequences of Claim 10 having at least a 10-base homopurine or homopyrimidine sequence, said probe comprising single-stranded DNA having at least a 10-base homopurine or homopyrimidine sequence and being adapted to bind to the major groove of double stranded DNA which includes said polynucleotide-encoding sequence.
- 25. The polynucleotide of Claim 1, wherein said SEQ ID NO is 913.
- 15 26. The polynucleotide of Claim 1, wherein said SEQ ID NO is 1039.
 - 27. The polynucleotide of Claim 1, wherein said SEQ ID NO is 1395.
- 28. The polynucleotide of Claim 1, wherein said SEQ ID NO is 1567.
 - 29. The polynucleotide of Claim 1, wherein said SEQ ID NO is 1667.
 - 30. The polynucleotide of Claim 1, wherein said SEQ ID NO is 1704.
- 25 31. The polynucleotide of Claim 1, wherein said SEQ ID NO is 2089.
 - 32. The polynucleotide of Claim 1, wherein said SEQ ID NO is 2297.
- 33. The polynucleotide of Claim 1, wherein said SEQ ID NO is 2302.

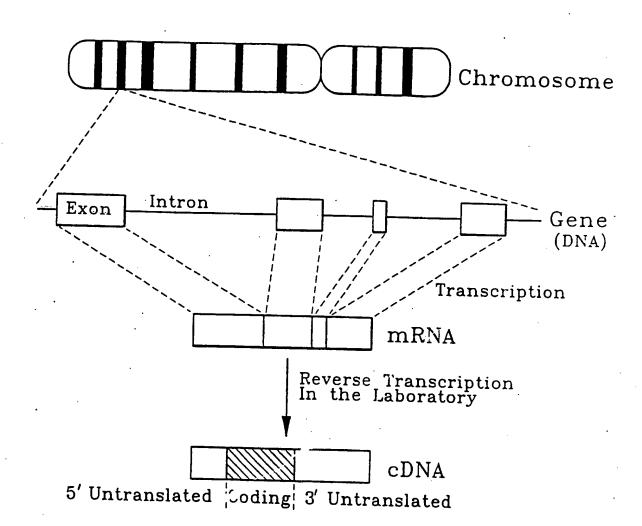


FIG. 1